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(54) Title: USE OF NEUREGULINS AS MODULATORS OF CELLULAR COMMUNICATION

(57) Abstract

The present invention relates to methods of affecting cellular communication in a vertebrate. The communication is affected by the administration of a neuregulin to a vertebrate, where the neuregulin interacts with a first cell type which results in the production of a product (i.e., Product A). This product, in turn, affects the function of a second cell type. Methods are disclosed in which the affect in function of the second cell type, results in the production of a second product (i.e., Product B) which, in turn, can affect the function of the first cell type or a third cell type. Additional methods are included for treatment of disorders involving an altered or inadequate level of production of a product involved in cellular communication.

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USE OF NEUREGULINS AS MODULATORS OF CELLULAR COMMUNICATION

5 FIELD OF THE INVENTION

This invention relates to methods of affecting cellular communication.

BACKGROUND OF THE INVENTION

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Vertebrate cells depend on externally produced factors for growth, differentiation and survival. These factors can be in the form of diffusible molecules that act at a distance from their site of synthesis. Alternatively these factors can be in the form of cell-surface-bound molecules that rely on cell-to-cell contact for their function. In many cases, different cell types may interact in a reciprocal manner in that both cell types produce factors that affect the other cell type. Vertebrates rely on these reciprocal interactions during embryogenesis and during the response to injury and disease.

Interdependence of cells and tissues plays important roles in the vertebrate nervous system. The nervous system is composed of neurons and neuroglial support cells. Peripheral nervous system axons are ensheathed by neuroglial cells (Schwann cells) and target organs which include skin, sensory receptors, muscle and other neurons. Additionally, peripheral axons interact with components of the central nervous system in the spinal cord. These include neurons and neuroglial cells such as astrocytes and oligodendrocytes.

It is well established that neurons and the tissues and cells with which they interact are dependent on each other for trophic support. This relationship is mediated by factors (proteins) produced by neurons that maintain the viability of target tissues (e.g. motor neuron derived factors that maintain muscle integrity) and neurotrophic factors produced by target (and other) tissues that maintain neuronal viability (e.g. muscle derived factors that maintain motor neuron viability). This interdependence plays an important role in embryonic development, maintenance of viability and response to injury in the nervous system and its targets.

The survival of various neuronal populations has been thought to be dependent only upon neurotrophic factors produced by targets of innervation. Recently it has been realized that neurotrophic factors are also derived from axonally associated cells

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(periaxonal glia), soma associated (perisomatic) cells (e.g. glia and efferent synapses) and from autocrine sources. These proteins are taken up by neurons where they exert their effect at the cell body. Neurotrophic factors either maintain the viability of the neuron or induce specific effects such as axonal extension, sprouting and other responses to injury and disease. Examples include factors such as nerve growth factor (NGF), brain derived neurotrophic factor (BDNF) and related molecules as well as ciliary neurotrophic factor (CNTF), insulin like growth factor (IGF) and fibroblast growth factors (FGF's) that all have neurotrophic activity and are derived from neuronally associated tissues as diverse as muscle, Schwann cells and spinal cord astrocytes and other neurons (e.g., Nishi, Science (1994) 265:1052).

The identification of pharmaceutical products or agents which induce the endogenous production of trophic factors would be beneficial treatment of diseases which involve trophic support.

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SUMMARY OF THE INVENTION

In general, the present invention relates to methods of affecting cellular communication in a vertebrate. The communication is affected by the administration of a neuregulin to a vertebrate, where the neuregulin interacts with a first cell type which results in the production of a product or products (i.e., Product(s) A). This product, in turn, affects the function of a second cell type (see Figures 9 and 10).

Neuregulins are a family of protein factors encoded by one gene. A variety of messenger RNA splicing variants (and their resultant proteins) are derived from this gene and many of these products show binding and activation to erbB2 (neu) and closely related receptors erbB3 and erbB4. The invention provides methods for using all of the known products of the neuregulin gene, as well as, other not yet discovered splicing variants of the neuregulin gene.

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Methods also are provided by the invention in which the effect in function of the second cell type, as described above, results in the production of a second product (i.e., Product B) which, in turn, can affect the function of the first cell type or a third cell type (see Figures 9 and 10).

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Included in the invention as well, are methods for treatment when disorders involve an altered or inadequate level of production of a product involved in cellular communication.

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Advantages of the present invention include the development of new therapeutic approaches to injury or disease based on the interdependence or communication of cells and the ability to influence or affect that communication with neuregulins. For example, a neuregulin factor that is produced by the second cell type can induce the first cell type to produce a product or products (Product(s) A) that are trophic for the second cell type. More specifically, cells and tissues that are associated with neurons may be induced to respond to a neuronally produced factor (neuregulin). This response would be in the form of the production of products (Product(s) A) that are trophic for neurons. The endogenous induction of more than one neurotrophic products by the neuregulin would be more effective than the therapeutic use of a single neurotrophic factor. Neurotrophic factors generally have restricted effects on specific neuronal subtypes (e.g. CNTF is trophic for motor neurons and NGF is trophic for sympathetic neurons as well as a subset of sensory neurons). Furthermore, the types of neurotrophic factors produced by a particular tissue are probably dependent on the target neuron type as well

as the type and stage of injury. As an example, CNTF, which is trophic for motor neurons, is released by Schwann cells in the early stages of recovery from nerve injury. This is replaced within a few days by Schwann cell and muscle derived BDNF, another motor neuron trophic factor (Curtis, et al., Nature (1993) 365:253-255; and Funakoshi, et al, J. Cell Biol. (1993) 123:455-465). In addition multiple neurotrophic factors function in vivo and may be synergistic in their effects. For example, it has been shown that multiple factors more efficiently arrest disease induced neuronal degeneration in animals than the use of a single factor (Mitsumoto et al., Science (1994) 265:1107).

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In the central nervous system, the neuregulin target, the first cell type, could be a neuron that in turn produces Product(s) A. Product A then affects other tissues (the second cell type) that produce neurotrophic products (Product(s) B) that affect the second cell type (the second cell type may be the source of the neuregulin), or perhaps a third cell type.

Thus, the use of the neuregulins, that are trophic for neuronally associated tissues in the manner described above would be therapeutically useful. Treatment would ensure the production of target specific combinations of products that are tailored to a particular disease state.

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BRIEF DESCRIPTION OF THE FIGURES

Figure 1 is a schematic diagram showing the method used to set up the SCG (superior cervical ganglion)/culture tube experiments; (1) tubes are filled with collagen +/- GGF; (2) SCG explants are placed in tubes; (3) tubes are cultured in humidified chambers; and (4) the extruded gels are fixed and stained as a "whole mount" (Anti-S100 for Schwann cells, and anti-tubulin β3 for axons.).

Figure 2 is a schematic diagram of the grid reticule inserted in the microscope ocular, which at a total magnification of 160X, allowed quantification of Schwann cell outgrowth and neurite outgrowth for the SCG/culture tube experiments.

Figure 3A shows the control data, that is, Schwann cell number as a function of distance from the SCG explant, for the SCG/culture tube experiments.

Figure 3B shows experimental data, of Schwann cell outgrowth for the SCG/culture tube experiments, at a dosage of 5 ng/ml rhGGF2.

Figure 3C shows experimental data, of Schwann cell outgrowth for the SCG/culture tube experiments, at a dosage of 50 ng/ml rhGGF2.

Figure 3D shows experimental data, of Schwann cell outgrowth for the SCG/culture tube experiments, at a dosage of 500 ng/ml rhGGF2.

Figure 4 shows the total number of Schwann cells as a function of days *in vitro* for the SCG/culture tube experiments.

Figure 5 shows experimental data, of neurite outgrowth, as a function of distance from the SCG explant, for the SCG/culture tube experiments performed at dosage levels of 5, 50 and 500 ng/ml rhGGF2.

Figure 6A shows a 2-dimensional dose-response matrix, used to examine the effects of rhGGF2 on neuronal survival and outgrowth.

Figure 6B illustrates the manner of counting, used in the afore-mentioned 2-dimensional dose-response experiment, by showing a representative sample well with fields of view.

Figure 7 shows experimental data of the effects of rhGGF2 on neuronal survival and outgrowth.

- Figure 8A shows data on the effects of exogenous GGF on the number of myelinated axons at 28 days post-injury.
 - Figure 8B shows the above-referenced data in bar graph form.
- Figure 9 represents a schematic illustration of the effect neuregulins can have on cellular communication.
 - Figure 10 represents a schematic illustration of specific effects of neuregulins on cellular communication within the nervous system.
- Figure 11A is a listing of the coding strand DNA sequence and deduced amino aid sequence of the cDNA obtained from splicing pattern of GGF2BPP1 shown in Figure 12. Potential glycosylation sites are underlined (along with polyadenylation signal AATAAA);
- Figure 11B is a listing of the coding strand DNA sequence and deduced amino acid sequence of the cDNA obtained from splicing pattern of GGF2BPP2. Potential glycosylation sites are underlined (along with polyadenylation signal AATAAA);
- Figure 11C is a listing of the coding strand DNA sequence and deduced amino acid sequence of the cDNA obtained from splicing pattern of GGF2BPP3. Potential glycosylation sites are underlined (along with polyadenylation signal AATAAA).
- Figure 12 is a diagram of representative splicing variants corresponding to bovine GGF gene products. The coding segments are represented by F, E, B, A, G, C, C/D, C/D', D, D', H, K and L. The location of the peptide sequences derived from purified protein are indicated by "o."
 - Figure 13 is a listing of the DNA sequences and predicted peptide sequences of the coding segments of GGF. Line 1 is a listing of the predicted amino acid sequences of bovine GGF, line 2 is a listing of the nucleotide sequences of bovine GGF, line 3 is a listing of the nucleotide sequences of human GGF (heregulin) (nucleotide base matches are indicated with a vertical line) and line 4 is a listing of the predicted amino acid sequences of human GGF/heregulin where it differs from the predicted bovine

sequence. Coding segments E, A' and K represent only the bovine sequences. Coding segment D' represents only the human (heregulin) sequence.

Figure 14 is the predicted GGF2 amino acid sequence and nucleotide sequence of BPP5. The upper line is the nucleotide sequence and the lower line is the predicted amino acid sequence.

Figure 15 is the predicted amino acid sequence and nucleotide sequence of GGF2BPP2. The upper line is the nucleotide sequence and the lower line is the predicted amino acid sequence.

Figure 16 is the predicted amino acid sequence and nucleotide sequence of GGF2BPP4. The upper line is the nucleotide sequence and the lower line is the predicted amino acid sequence.

Figure 17 is a list of splicing variants derived from the sequences shown in Figure 13.

Figure 18 is the predicted amino acid sequence, bottom, and nucleic sequence, top, of EGFL1.

Figure 19 is the predicted amino acid sequence, bottom, and nucleic sequence, top, of EGFL2.

Figure 20 is the predicted amino acid sequence, bottom, and nucleic sequence, top, of EGFL3.

Figure 21 is the predicted amino acid sequence, bottom, and nucleic sequence, top, of EGFL4.

Figure 22 is the predicted amino acid sequence, bottom, and nucleic sequence, top, of EGFL5.

Figure 23 is the predicted amino acid sequence, bottom, and nucleic sequence, top, of EGFL6.

Figure 24 is the predicted amino acid sequence (middle) and nucleic sequence (top) of GGF2HBS5. The bottom (intermittent) sequence represents peptide sequences derived from GGF-II preparations.

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Figure 25 is the sequences of GGFHBS5, GGFHB1 and GGFBPP5 polypeptides.

Figure 26 is the amino acid sequence of cDNA encoding mature hGGF2.

Figure 27 depicts a stretch of the putative bovine GGF-II gene sequence from the recombinant bovine genomic phage GGF2BG1. The figure is the coding strand of the DNA sequence and the deduced amino acid sequence in the third reading frame.

DETAILED DESCRIPTION OF THE INVENTION

It is intended that all references cited shall be incorporated herein by reference.

5 General

The invention pertains to methods of affecting cellular communication in vertebrates. The communication is affected by the administration of a neuregulin to a vertebrate where the neuregulin interacts with a first cell type which results in the production of a product. This product, in turn, affects the function of a second cell type. More specifically, the invention relates to the induction of endogenous tropic factors (products) by the administration of a neuregulin.

Methods also are provided by the invention in which the affect in function of the second cell type, described above, results in the production of a second product which, in turn, can affect the function of the first cell type, the second cell type or a third cell type.

Definition of Key Terms

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The term <u>administration</u> as used herein refers to the act of delivering a substance, including but not limited to the following routes: parenteral, intravenous, subcutaneous, intramuscular, intraorbital, ophthalmic, intraventricular, intracranial, intracapsular, intraspinal, intracisternal, intraperitoneal, topical, intranasal, aerosol, scarification, orally, buccal, rectal or vaginal. Administration as used herein refers to a pharmaceutical preparation of a substance and the delivery of that preparation to a recipient.

The term <u>affecting</u> as used herein refers to the induction of a quantitative change in the response of a target cell, as a result of an interaction with a Product.

The term <u>Alzheimer's Disease</u> as used herein refers to a progressive central neurodegeneration involving loss of cortical and other neurons, and associated with neurofibrillary tangles and \(\mathbb{B}\)-amyloid deposits.

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The term <u>amyotrophic lateral sclerosis</u> (ALS) as used herein refers to a motor neuron disease characterized by a progressive degeneration of the neurons that give rise to the corticospinal tract that results in a deficit in upper and lower motor neurons.

The term <u>astrocyte</u> as used herein refers to a neuroglial cell of ectodermal origin and its progenitor cells. This cell has a round nucleus and a "star shaped" body and many long processes that end as vascular foot plates on the small blood vessels of the CNS and is associated with other structures. A more complete definition of the astrocyte and its progenitors can be found in the following materials: Reynolds and Weiss, *Science* (1992) 255:1707-1710; Reynolds, Tetzlaff, and Weiss, *J. Neurosci* (1992) 12:4565-4574; and Kandel, et al., *Principles of Neuroscience*, third ed. (1991), Appleton & Lange, Norwalk, CT.

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The term <u>cellular communication</u> as used herein refers to the synthesis of a substance in one cell type and the interaction of that substance with a second cell type. This process includes, but is not limited to, secretion of the substance from a cell. The substance elicits a change in the second cell type or with the first cell type. Communication can occur reciprocally or non-reciprocally with one or more cell types.

The term <u>differentiation</u> as used herein refers to a morphological and/or chemical change that results in the generation of a different cell type or state of specialization. The differentiation of cells as used herein refers to the induction of a cellular developmental program which specifies one or more components of a cell type. The therapeutic usefulness of differentiation can be seen, in increases in quantity of any component of a cell type in diseased tissue by at least 10% or more, more preferably by 50% or more, and most preferably by more than 100% relative to the equivalent tissue in a similarly treated control animal.

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The term <u>disorder</u> as used herein refers to a disturbance of function and/or structure of a living organism, resulting from an external source, a genetic predisposition, a physical or chemical trauma, or a combination of the above, including but not limited to any mammalian disease.

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The term <u>first cell type</u> as used herein refers to the cell type that interacts with a neuregulin. The first cell type includes but is not limited to one or more of the following: neuron, glial cell, Schwann cell, astrocyte, oligodendrocyte, myoblast, muscle cell, satellite cell, skin cell, sensory organ cell, inflammatory cell such as macrophage, neutrophil, T-cell, eosinophil, mast cell, basophil and stromal cell such as fibroblasts or endothelial cells. Bloom and Fawcett, *A Textbook of Histology*, tenth ed. (1975), W. B. Saunders Company, Philadelphia, PA.

The term <u>function</u> as used herein refers to any activity or response of a cell. These include but are not limited to proliferation, differentiation, growth, survival, changes in the pattern of gene expression and secretion, and metabolic changes.

The term <u>glial cell</u> as used herein refers to connective and support tissues of the nervous system and includes ectodermally derived astrocytes, oligodendroglia, Schwann cells and mesodermally derived microglia and their progenitors. A more complete definition of glial cells and their progenitors can be found in the following materials: Anderson, *FASEB J.* (1994) 8:707-713; Reynolds and Weiss, *Science* (1992) 255:1707-1710; Reynolds, Tetzlaff, and Weiss, *J. Neurosci* (1992) 12:4565-4574; and Kandel, et al., *Principles of Neuroscience*, third ed. (1991), Appleton & Lange, Norwalk, CT.

The term <u>interacts</u> as used herein refers to a contact with a target (cell), including but not limited to binding of a product to a cell receptor.

The term <u>mammal</u> as used herein describes a member of the Class Mammalia (Subphylum Vertebrata).

The term <u>matrix molecule</u> as used herein refers to a chemical component of the insoluble meshwork of extracellular proteins that mediate adhesive interactions between cells and modulate the functions of cells.

The term <u>mitosis</u> as used herein refers to the division of a cell where each daughter nucleus receives identical complements of the numbers of chromosomes characteristic of the somatic cells of the species. Mitosis as used herein refers to any cell division which results in the production of new cells in the patient. More specifically, a useful therapeutic is defined *in vitro* as an increase in mitotic index relative to untreated cells of 50%, more preferably 100%, and most preferably 300%, when the cells are exposed to labeling agent for a time equivalent to two doubling times. The mitotic index is the fraction of cells in the culture which have labeled nuclei when grown in the presence of a tracer which only incorporates during S phase (i.e., BrdU) and the doubling time is defined as the average time required for the number of cells in the culture to increase by a factor of two.

For example, one effect on mitosis in vivo is defined as an increase in satellite cell activation as measured by the appearance of labeled satellite cells in the muscle tissue of a mammal exposed to a tracer which only incorporates during S phase (i.e.,

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BrdU). A useful therapeutic is defined in vivo as a compound which increases satellite cell activation relative to a control mammal by at least 10%, more preferably by at least 50%, and most preferably by more than 200% when the mammal is exposed to labeling agent for a period of greater than 15 minutes and tissues are assayed between 10 hours and 24 hours after administration of the mitogen at the therapeutic dose.

The term <u>muscle cell</u> as used herein refers to a cellular component of skeletal, smooth or cardiac muscle, including but not limited to myofibrils, satellite cells, and myoepithelial cells and their progenitors. A more complete definition of muscle cells can be found in, Wheater, et al., *Functional Histology* (1987), Churchill Livingstone, New York, NY; and *Myology*, ed. by Engel and Franzini-Armstrong, second ed. (1994) McGraw Hill, New York, NY.

The term neuregulin as used herein refers to the glial growth factors, the heregulins, neu differentiation factor, acetylcholine receptor inducing activity, and erbB2, 3 and 4 binding proteins. A more complete definition of neuregulins can be found in the specification herein and in the following materials: U.S. Patent No. 5,237,056; U.S. Patent Application SN 08/249,322; WO 92/20798; EPO 0 505 148 A1; Marchionni, et al., Nature 362:313, 1993; Benveniste, et al., PNAS 82:3930-3934, 1985; Kimura, et al., Nature (1990) 348:257-260; Davis and Stroobant, J. Cell. Biol. (1990) 110:1353-1360; Wen, et al., Cell (1992) 69:559; Yarden and Ullrich, Ann. Rev. Biochem. (1988) 57:443, ; Holmes, et al., Science 256:1205, 1992; Dobashi, et al., Proc. Nat'l. Acad. Sci. 88:8582, 1991; Lupu, et al., Proc. Nat'l. Acad. Sci. (1992) 89:2287; Peles and Yarden, BioEssays (1993) 15:815, Mudge, Current Biology (1993) 3:361, all hereby incorporated by reference.

The term <u>neuregulin producing cell</u> as used herein refers to a cell that produces a neuregulin. The term refers to all producer cells including cells that produce recombinant neuregulins.

The term <u>neurological disorder</u> as described herein refers to a disorder of the nervous system.

The term <u>nervous system cell</u> as used herein includes nervous system support cells and neurons.

The term <u>neuron</u> as used herein refers to a complete nerve cell, including the cell body and all of its processes, and its progenitors. A more complete definition of neuron

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and its progenitors can be found in the following materials: Reynolds and Weiss, Science (1992) 255:1707-1710; Reynolds, Tetzlaff, and Weiss, J. Neurosci (1992) 12:4565-4574; Ray, Peterson, Schinstine, and Gage, PNAS (1993) 90:3602-3606; and Kandel, et al., Principles of Neuroscience, third ed. (1991), Appleton & Lange, Norwalk, CT.

The term <u>neurotrophic agent</u> as used herein refers to a substance that elicits a trophic effect in one or more neuronal subtypes. These effects include but are not limited to survival, sprouting and differentiation.

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The term <u>oligodendrocyte</u> as used herein refers to the neuroglial cells, of ectodermal origin, with small oval nuclei and fine cytoplasmic processes that are responsible for the formation of myelin in the CNS. The progenitors of oligodendrocytes are also included. A more complete definition of oligodendrocytes and their progenitors can be found in Kandel, et al., *Principles of Neuroscience*, third ed. (1991), Appleton & Lange, Norwalk, CT.

The term <u>Parkinson's Disease</u> as used herein refers to a progressive central neurodegeneration involving dopaminergic neurons.

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The term <u>peripheral neuropathy</u> as used herein refers to functional disturbances and/or pathological changes in the peripheral nervous system.

The term <u>Product</u> as used herein refers to any substance as defined herein as Product A or Product B.

The term <u>Product A</u> as used herein refers to the substances whose synthesis and release are induced in the first cell type by neuregulin. Such substances include but are not limited to one or more of the following: nerve growth factor (NGF), neurotrophins, brain-derived neurotrophic factor, ciliary neurotrophic factor, leukemia inhibiting factor, interleukin 6, platelet derived growth factor, fibroblast growth factors, transforming growth factor B, epidermal growth factor, transforming growth factor a, neuregulins, insulin like growth factor, matrix molecules, adhesion molecules, growth factor receptors, low affinity NGF receptor, proteases, protease inhibitors, and antioxidants.

The term <u>Product B</u> as used herein refers to the substances whose synthesis and release are induced in the second cell type by Product A. Such substances include but

are not limited to one or more of the following: nerve growth factor (NGF), neurotrophins, brain-derived neurotrophic factor, ciliary neurotrophic factor, leukemia inhibiting factor, interleukin 6, platelet derived growth factor, fibroblast growth factors, transforming growth factor β, epidermal growth factor, transforming growth factor a, neuregulins, glial derived neurotrophic factor, insulin like growth factor, matrix molecules, adhesion molecules, growth factor receptors, low affinity NGF receptor (p75), proteases, protease inhibitors and antioxidants.

The term <u>production</u> as used herein refers to induced or constitutive synthesis and/or release of a Product from a cell.

The term <u>protease</u> as used herein refers to an enzyme that hydrolyses peptide bonds in a protein molecule.

The term <u>protease inhibitor</u> as used herein refers to a molecule that inhibits the activity and/or function of a protease.

The term <u>Schwann cell</u> as used herein refers to the neuroglial cell composing the neurolemma of peripheral nerve fibers and its progenitors. A more complete definition of Schwann cells and their progenitors can be found in the following materials: Anderson, *FASEB J.* (1994) 8:707-713; Kandel, et al., *Principles of Neuroscience*, third ed. (1991), Appleton & Lange, Norwalk, CT.

The term second cell type as used herein refers to the cell type that interacts with and responds to Product A. The second cell type includes but is not limited to one or more of the following: neuron, glial cell, Schwann cell, astrocyte, oligodendrocyte, myoblast, muscle cell, satellite cell, skin cell, sensory organ cell, inflammatory cell such as macrophage, neutrophil, T-cell, eosinophil, mast cell, basophil and stromal cell such as fibroblasts or endothelial cells. A more complete definition may be found in Bloom and Fawcett, A Textbook of Histology, tenth ed. (1975), W. B. Saunders Company, Philadelphia, PA.

The term <u>sensory organ cell</u> as used herein refers to a primary sensory cell contained within a sensory organ and its progenitors and includes but is not limited to one or more of the following: taste cells, olfactory epithelial cell, rod and cone photoreceptors, Meisner corpuscle, Ruffini corpuscle, Merckel receptor, Pacinian corpuscle, muscle spindle cell, cochleovestibular hair cells and joint mechanoreceptor cells. A more complete definition of sensory organ cells and their progenitors can be

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found in, Wheater, et al., Functional Histology (1987), Churchill Livingstone, New York, NY; Mahanthappa and Schwarting, Neuron (1993) 10:293-305; Forge, Li, Corwin and Nevill, Science (1993) 259:1616-1622; Tsue, Watling, Weisleder, Coltrera and Rubel, J. Neurosci (1994) 14:140-152.

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The term skin cell as used herein refers to the cellular components of the skin and includes fibroblasts, keratinocytes, epidermal cells, hair follicle cells, melanocytes, myoepithelial sweat gland cells, and sebaceous gland cells and their progenitors. A more complete definition of skin cells and their progenitors can be found in, Wheater, et al., Functional Histology (1987), Churchill Livingstone, New York, NY.

The term <u>spinal muscular atrophy</u> as used herein refers to a progressive disease of upper and lower motor neurons, usually present in childhood.

The term <u>survival</u> as used herein refers to any process where a cell avoids death. The term survival as used herein also refers to the prevention of cell loss as evidenced by necrosis or apoptosis or the prevention of other mechanisms of cell loss. Survival as used herein indicates a decrease in the rate of cell death of at least 10%, more preferably by at least 50%, and most preferably by the least 300% relative to an untreated control.

The rate of survival may be measured by counting cells stainable with a dye specific for dead cells (such as propidium iodide) in culture when the cells are 8 days post-differentiation (i.e., 8 days after the media is changed from 20% to 0.5% serum).

The term therapeutically effective amount as used herein refers to that amount which will produce a desirable result upon administration and which will vary depending upon a number of issues, including the dosage to be administered, and the route of administration.

The term third cell type as used herein refers to a cell type that interacts with and responds to Product B. The third cell type may be identical to the first cell type. The third cell type includes but is not limited to one or more of the following: neuron, glial cell, Schwann cell, astrocyte, oligodendrocyte, myoblast, muscle cell, satellite cell, skin cell, sensory organ cell, inflammatory cell such as macrophage, neutrophil, T-cell, eosinophil, mast cell, basophil and stromal cell such as fibroblasts or endothelial cells. A more complete definition may be found in Bloom and Fawcett, A Textbook of Histology, tenth ed. (1975), W. B. Saunders Company, Philadelphia, PA.

The term <u>treating</u> as used herein may refer to a procedure (e.g. medical procedure) designed to exert a beneficial effect on a disorder. Treating as used herein means any administration of a substance described herein for the purpose of increasing cellular communication of products. Most preferably, the treating is for the purpose of reducing or diminishing the symptoms or progression of a disease or disorder of cells. Treating as used herein also means the administration of a substance to increase or alter the cells in healthy individuals. The treating may be brought about by the contacting of the cells which are sensitive or responsive to the neuregulins described herein with an effective amount of the neuregulin.

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The term <u>trophic</u> as used herein refers to an effect of a substance on a cell, including but not limited to proliferation, growth, sprouting, differentiation or survival.

The term <u>vertebrate</u> as used herein refers to an animal that is a member of the Subphylum Vertebrata (Phylum Chordata).

Neuregulins

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A novel aspect of the present invention relates to the ability of neuregulins to affect cellular communication between different and similar cell types. Neuregulins are the products of a gene which produce a number of variably-sized, differentially-spliced RNA transcripts that give rise to a series of proteins. These proteins are of different lengths and contain some common peptide sequences and some unique peptide sequences. The conclusion that these factors are encoded by a single gene is supported by the differentially-spliced RNA sequences which are recoverable from bovine posterior pituitary, human spinal chord and human breast cancer cells (MDA-MB-231). Further support for this conclusion derives from the size range of proteins which act as ligands for the p185erbB2 receptor (see below).

Further evidence to support the fact that the genes encoding GGF/p185erbB2 binding proteins are homologous comes from nucleotide sequence comparison. Holmes et al., (Science (1992) 256:1205-1210) demonstrate the purification of a 45-kilodalton human protein (Heregulin-α) which specifically interacts with the receptor protein p185erbB2. Peles et al., (Cell (1992) 69:559) describe a complementary DNA isolated from rat cells encoding a protein call "neu differentiation factor" (NDF). The translation product of the NDF cDNA has p185erbB binding activity. Several other groups have reported the purification of proteins of various molecular weights with p185erbB2 binding activity. These groups include the following: Lupu et al., (1992) Proc. Nat'l. Acad. Sci. USA 89:2287; Yarden and Peles, (1991) Biochemistry 30:3543; Lupu et al., (1990) Science 249:1552; Dobashi et al., (1991) Biochem. Biophys. Res. Comm. 179:1536; and Huang et al., (1992) J. Biol. Chem. 257:11508-11512.

We have found that p185erbB2 and related receptor binding proteins (i.e., p185erbB3 and p185erbB4) affect cellular communication. This effect results in the production of a product from a first cell type, where the product, in turn affects the function of a second cell type. The affect in a function of the second cell type and can result in the production of other products which also can affect functions of other cell types. For example, neuregulins can interact with Schwann cells, which as a result of this interaction produce neurotrophic agents. These agents, in turn, interact with neurons to promote their neuronal regeneration. Alternatively, in the central nervous system, a first cell type, being a neuron, could produce a neuregulin, which in turn, affects a second cell type which is a neuron also.

These neuregulins may be identified using the protocols described herein (Examples 1 and 2) and in Holmes et al., Science (1992) 256: 1205; Peles et al., Cell (1992) 69:205; Wen et al., Cell (1992) 69:559; Lupu et al., Proc. Nat'l. Acad. Sci. USA (1992) 89:2287; Yarden and Peles, Biochemistry (1991) 30:3543; Lupu et al., Science (1990) 249:1552; Dobashi et al., Biochem. Biophys. Res. Comm. (1991) 179:1536; Huang et al., J. Biol. Chem. (1992) 257:11508-11512; Marchionni et al., Nature (1993) 362:313; and in U.S. Patent Application Serial No. 07/931,041, filed August 17, 1992, all of which are incorporated herein by reference.

Specifically, the invention provides for use of polypeptides of a specified formula, and DNA sequences encoding those polypeptides. The polypeptides have the formula

WYBAZCX

wherein WYBAZCX is composed of the amino acid sequences shown in Figure 13; wherein W comprises the polypeptide segment F, or is absent; wherein Y comprises the polypeptide segment E, or is absent; wherein Z comprises the polypeptide segment G or is absent; and wherein X comprises the polypeptide segments C/D HKL, C/D H, C/D HL, C/D D, C/D' HKL, C/D H, C/D H, C/D D, C/D HKL, C/D C/D' HKL, C/D C/D' H, C/D C/D' HKL, C/D C/D' D, C/D D' HKL, provided that, either

- a) at least one of F, Y, B, A, Z, C, or X is of bovine origin; or
- b) Y comprises the polypeptide segment E; or
- c) X comprises the polypeptide segments C/D HKL, C/D D, C/D' HKL, C/D C/D' HKL, C/D D' H, C/D D' H, C/D D' HKL, C/D D' HKL, C/D' D' H, C/D' D' HKL, C/D C/D' D' HKL, C/D C/D' D' HKL, C/D C/D' HKL, C/D C/D' HKL, C/D C/D' HL.

In addition, the invention includes the use of the DNA sequence comprising coding segments ⁵FBA³ as well as the corresponding polypeptide segments having the amino acid sequences shown in Figure 13;

the DNA sequence comprising the coding segments ⁵FBA'³ as well as the corresponding polypeptide segments having the amino acid sequences shown in Figure 13;

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the DNA sequence comprising the coding segments ⁵FEBA³ as well as the corresponding polypeptide segments having the amino acid sequences shown in Figure 13;

the DNA sequence comprising the coding segments ⁵FEBA'³ as well as the corresponding polypeptide segments having the amino acid sequences shown in Figure 13;

the DNA sequence comprising the polypeptide coding segments of the GGF2HBS5 cDNA clone (ATCC Deposit No. 75298, deposited September 2, 1992), also known as GGF-II.

The invention further includes the use of peptides of the formula FBA, FEBA, FBA' FEBA' and DNA sequences encoding these peptides wherein the polypeptide segments correspond to amino acid sequences shown in Figure 13. The purified GGF-II polypeptide is also included as part of the invention.

Also included in this invention is the mature GGF peptide and the DNA encoding said peptide, exclusive of the N-terminal signal sequence, which is also useful for treatment of conditions involving abnormalities in cellular communication.

Furthermore, the invention includes a method of cellular communication by the application to a vertebrate of a

- 30 kD polypeptide factor isolated from the MDA MB 231 human breast cell line; or
 - 35 kD polypeptide factor isolated from the rat I-EJ transformed fibroblast cell line to the glial cell; or
 - -75 kD polypeptide factor isolated from the SKBR-3 human breast cell line; or
- -44 kD polypeptide factor isolated from the rat I-EJ transformed fibroblast cell 30 line, or
 - -25kD polypeptide factor isolated from activated mouse peritoneal macrophages; or
 - -45 kD polypeptide factor isolated from the MDA MB 231 human breast cell; or
- -7 to 14 kD polypeptide factor isolated from the ATL-2 human T-cell line to the glial cell; or
 - -25 kD polypeptide factor isolated from the bovine kidney cell; or
 - -42 kD polypeptide factor (ARIA) isolated from brains.

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The invention further includes a method for the use of the EGFL1, EGFL2, EGFL3, EGFL4, EGFL5 and EGFL6 polypeptides, Figure 18 to Figure 23 respectively, and for the methods of affecting cellular communication in vivo and in vitro.

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Also included in the invention is the administration of the GGF-II polypeptide whose sequence is shown in Figure 24, for affecting cellular communication.

An additional aspect of the invention includes the use of the above-referenced peptides for the purpose of stimulating Schwann cells to produce growth factors which may, in turn, be harvested for scientific or therapeutic use.

Thus, the invention further embraces a polypeptide factor capable of affecting cellular communication and including an amino acid sequence encoded by:

- (a) a DNA sequence shown in Figure 11;
- (b) a DNA sequence shown in Figure 27;
- (c) the DNA sequence represented by nucleotides 281-557 of the sequences shown in Figure 11; or
- (d) a DNA sequence hybridizable to any one of the DNA sequences 20 according to (a), (b) or (c).

The invention further includes sequences which have greater than 60%, preferably 80%, sequence identity of homology to the sequences indicated above.

While the present invention is not limited to a particular set of hybridization conditions, the following protocol gives general guidance which may, if desired, be followed:

DNA probes may be labeled to high specific activity (approximately 10⁸ to 10⁹ 32 Pdmp/μg) by nick-translation or by PCR reactions according to Schowalter and Sommer (Anal. Biochem. (1989) 177:90-94) and purified by desalting on G-150 Sephadex columns. Probes may be denatured (10 minutes in boiling water followed by immersion into ice water), then added to hybridization solutions of 80% buffer B (2g polyvinylpyrolidine, 2g Ficoll-400, 2g bovine serum albumin, 50ml 1M Tris HCl (pH 7.5), 58g NaCl, 1g sodium pyrophosphate, 10g sodium dodecyl sulfate, 950 ml H₂O) containing 10% dextran sulfate at 10⁶ dpm ³²P per ml and incubated overnight (approximately 16 hours) at 60° C. The filters may then be washed at 60° C first in

buffer B for 15 minutes followed by three 20-minute washes in 2X SSC, 0.1% SDS then one for 20 minutes in 1XSSC, 0.1% SDS.

In other respects, the invention provides:

5 (a) a basic polypeptide factor which has, if obtained from bovine pituitary material, an observed molecular weight, whether in reducing conditions or not, of from about 30kD to about 36kD on SDS-polyacrylamide gel electrophoresis using the following molecular weight standards:

	Lysozyme (hen egg white)	14,400
10	Soybean trypsin inhibitor	
	Carbonic anhydrase (bovine)	31,000
	Ovalbumin (hen egg white)	45,000
	Bovine serum albumin	
	Phosphorylase B (rabbit muscle)	97,400;

which factor has glial cell mitogenic activity including stimulating the division of rat Schwann cells in the presence of fetal calf plasma, and when isolated using reversed-phase HPLC retains at least 50% of said activity after 10 weeks incubation in 0.1 % trifluoroacetic acid at 4° C; and

20 (b) a basic polypeptide factor which has, if obtained from bovine pituitary material, an observed molecular weight, under non-reducing conditions, of from about 55 kD to about 63 kD on SDS-polyacrylamide gel electrophoresis using the following molecular weight standards:

	Lysozyme (hen egg white)	14,400
25	Soybean trypsin inhibitor	
	Carbonic anhydrase (bovine)	31,000
	Ovalbumin (hen egg white)	45,000
	Bovine serum albumin	
	Phosphorylase B (rabbit muscle)	97,400;

which factor the human equivalent of which is encoded by DNA clone GGF2HBS5 described herein and is capable of affecting cellular communication.

For convenience of description only, the lower molecular weight and higher molecular weight factors of this invention are referred to hereafter as "GGF-I" and "GGF-II", respectively. The "GGF2" designation is used for all clones isolated with peptide sequence data derived from GGF-II protein (i.e., GGF2HBS5, GGF2BPP3).

It will be appreciated that the molecular weight range limits quoted are not exact, but are subject to slight variations depending upon the source of the particular polypeptide factor. A variation of, say, about 10% would not, for example, be impossible for material from another source.

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Another important aspect of the invention is a DNA sequence encoding a polypeptide capable of affecting cellular communication and comprising:

- (a) a DNA sequence shown Figure 11;
- (b) a DNA sequence shown in Figure 27;
- (c) the DNA sequence represented by nucleotides 281-557 of the sequence shown in Figure 11; or
 - (d) a DNA sequence hybridizable to any one of the DNA sequences according to (a), (b) or (c).

Thus other important aspects of the invention are:

- (a) A series of human and bovine polypeptide factors capable of affecting cellular communication. These peptide sequences are shown in Figures 13, 14, 15 and 16, respectively.
- (b) A series of polypeptide factors capable of affecting cellular communication and purified and characterized according to the procedures outlined by Lupu et al., Science (1990) 249:1552; Lupu et al., Proc. Nat'l. Acad. Sci USA (1992) 89: 2287; Holmes et al., Science (1992) 256:1205; Peles et al., Cell (1992) 69:205; Yarden and Peles, Biochemistry (1991) 30:3543; Dobashi et al., Proc. Nat'l. Acad. Sci. (1991) 88: 8582; Davis et al., Biochem. Biophys. Res. Commun. (1991) 179:1536; Beaumont et al., Patent Application PCT/US91/03443 (1990); Greene et al., Patent Application PCT/US91/02331 (1990); Usdin and Fischbach, J. Cell. Biol. (1986) 103:493-507; Falls et al., Cold Spring Harbor Symp. Quant. Biol. (1990) 55:397-406; Harris et al., Proc. Nat'l. Acad. Sci. USA (1991) 88:7664-7668; and Falls et al., Cell (1993) 72:801-815.
 - (c) A polypeptide factor (GGFBPP5) capable of affecting cellular communication. The amino acid sequence is shown in Figure 14, and is encoded by the bovine DNA sequence shown in Figure 14.

The novel human peptide sequences described above and presented in Figures 13, 14, 15, and 16, respectively, represent a series of splicing variants which can be isolated as full length complementary DNAs (cDNAs) from natural sources (cDNA libraries prepared from the appropriate tissues) or can be assembled as DNA constructs with individual exons (e.g., derived as separate exons) by someone skilled in the art.

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Other compounds in particular, peptides, which bind specifically to the p185erbB2 receptor and related receptors can also be used according to the invention as affections of cellular communication. A candidate compound can be routinely screened for p185erbB2 binding, and, if it binds, it can then be screened for affecting cellular communication using the methods described herein.

The invention includes any modifications or equivalents of the above polypeptide factors which do not exhibit a significantly reduced activity. For example, modifications in which amino acid content or sequence is altered without substantially adversely affecting activity are included. By way of illustration, in EP-A 109748, mutations of native proteins are disclosed in which the possibility of unwanted disulfide bonding is avoided by replacing any cysteine in the native sequence which is not necessary for biological activity with a neutral amino acid. The statements of effect and use contained herein are therefore to be construed accordingly, with such uses and effects employing modified or equivalent factors being part of the invention.

The new sequences of the invention open up the benefits of recombinant technology. The invention thus also includes the following aspects:

- (a) DNA constructs comprising DNA sequences as defined above in operable reading frame position within vectors (positioned relative to control sequences so as to permit expression of the sequences) in chosen host cells after transformation thereof by the constructs (preferably the control sequence includes regulatable promoters, e.g. Trp). It will be appreciated that the selection of a promoter and regulatory sequences (if any) are matters of choice for those of skill in the art:
- (b) host cells modified by incorporating constructs as defined in (a) immediately above so that said DNA sequences may be expressed in said host cells the choice of host is not critical, and chosen cells may be prokaryotic or eukaryotic and may be genetically modified to incorporate said constructs by methods known in the art; and,
- (c) a process for the preparation of factors as defined above comprising cultivating the modified host cells under conditions permitting expression of the DNA sequences. These conditions can be readily determined, for any particular embodiment, by those of skill in the art of recombinant DNA technology. Glial cell mitogens prepared by this means are included in the present invention.

None of the factors described in the art has the combination of characteristics possessed by the present new polypeptide factors.

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The invention also includes a neuregulin as defined above, by extracting vertebrate brain material to obtain protein, subjecting the resulting extract to chromatographic purification by hydroxyapatite HPLC and then subjecting these fractions to SDS-polyacrylamide gel electrophoresis. The fraction which as an observed molecular weight of about 30kD to 36kD and/or the fraction which has an observed molecular weight of about 55kD to 63kD is collected. In either case, the fraction is subjected to SDS-polyacrylamide gel electrophoresis using the following molecular weight standards:

	Lysozyme (hen egg white)	14,400
10	Soybean trypsin inhibitor	21,500
	Carbonic anhydrase (bovine)	31,000
	Ovalbumin (hen egg white)	45,000
	Bovine serum albumin	66,200
	Phosphorylase B (rabbit muscle)	97,400

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In the case of the smaller molecular weight fraction, the SDS-polyacrylamide gel is run in non-reducing conditions or in reducing conditions and in the case of the larger molecular weight fraction, the gel is run under non-reducing conditions. The fractions are then tested for activity stimulating the division of rat Schwann cells against a background of fetal calf plasma.

Preferably, the above process starts by isolating a relevant fraction obtained by carboxymethyl cellulose chromatography, e.g. from bovine pituitary material. It is also preferred that hydroxyapatite HPLC, cation exchange chromatography, gel filtration, and/or reversed-phase HPLC be employed prior to the SDS-Polyacrylamide gel electrophoresis. At each stage in the process, activity may be determined using Schwann cell incorporation of radioactive iododeoxyuridine as a measure in an assay generally as described by Brockes in *Meth. Enz.* (1987) 147:217-225, but modified by substituting 10% FCP for 10% FCS. As already noted, such as assay is an aspect of the invention in its own substance for CNS or PNS cell, e.g. Schwann cell, mitogenic effects.

Compounds may be assayed for their usefulness in vitro using the methods provided in the description and examples below. Following the in vitro demonstration of the effect of neuregulins on cellular communication between various cell types, the in vivo therapeutic benefit of the effect can be accomplished by the administration of neuregulins, neuregulin producing cells or DNA encoding neuregulins to a vertebrate

requiring therapy. In a specific example, in vivo testing can be demonstrated as described in Example 3.

The invention includes the use of the above named family of proteins (i.e. neuregulins) as extracted from natural sources (tissues or cell lines) or as prepared by recombinant means.

Other compounds in particular, peptides, which bind specifically to the p185erbB2 and related receptor binding proteins (i.e., p185erbB3 and p185erbB4) can also be used according to the invention as affectors of cellular communication. A candidate compound can be routinely screened for p185erbB2, p185erbB3 and p185erbB4 binding, and if it binds, can then be screened for affecting cellular communication using the methods described herein.

The invention includes use of any modifications or equivalents of the above polypeptide factors which do not exhibit a significantly reduced activity related to affecting cellular communication. For example, modifications in which amino acid content or sequence is altered without substantially adversely affecting activity are included. The statements of effect and use contained herein are therefore to be construed accordingly, with such uses and effects employing modified or equivalent factors being part of the invention.

The human peptide sequences described above represent a series of splicing variants which can be isolated as full length complementary DNAs (cDNAS) from natural sources (cDNA libraries prepared from the appropriate tissues) or can be assembled as DNA constructs with individual exons (e.g., derived as separate exons) by someone skilled in the art.

The invention includes methods for the use of any protein which is substantially homologous to the coding segments in Figure 13, as well as other naturally occurring GGF polypeptides for the purpose of inducing muscle mitogenesis. Also included are the use of: allelic variations; natural mutants; induced mutants; proteins encoded by DNA that hybridizes under high or low stringency conditions to a nucleic acid naturally occurring (for definitions of high and low stringency see *Current Protocols in Molecular Biology*, (1989) John Wiley & Sons, New York, NY, 6.3.1 - 6.3.6, hereby incorporated by reference); and the use of polypeptides or proteins specifically bound by antisera to GGF polypeptides. The term also includes the use of chimeric polypeptides that include the GGF polypeptides comprising sequences from Figure 13.

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Use of Neuregulins

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A novel aspect of the invention involves the use of neuregulins as factors to promote cell communication by inducing the production of products. These Products affect the function of these cells. Treatment of the cells to achieve these effects may be achieved by contacting cells with a polypeptide described herein.

The methods of the invention may also be used to treat a patient suffering from a disease caused by a lack of trophic factor(s). The lack of trophic factor(s) is defined by a decreased amount of trophic factor(s) relative to that of an unaffected individual sufficient to cause detectable alteration in the biological effect of those trophic factor(s). The neurotrophic factor(s) may be present at levels 10% below those observed in unaffected individuals. More preferably, the factor(s) are present at levels 20% lower than that observed in unaffected individuals, and most preferably the levels are lowered by 80% relative to unaffected individuals under similar circumstances.

The methods of the invention make use of the fact that the neuregulin proteins are encoded by the same gene. A variety of messenger RNA splicing variants (and their resultant proteins) are derived from this gene and many of these products show binding to p185erbB2 and related receptors and activation of the same. This invention provides a use for all of the known products of the neuregulin gene (described herein and in the references listed above). Most preferably, recombinant human GGF2 (rhGGF2) is used in these methods.

The invention also relates to the use of other, not yet naturally isolated, splicing variants of the neuregulin gene. Figure 12 shows the known patterns of splicing. These patterns are derived from polymerase chain reaction experiments (on reverse transcribed RNA), analysis of cDNA clones (as presented within), and analysis of published sequences encoding neuregulins (Peles et al., Cell (1992) 69:205 and Wen et al., Cell (1992) 69:559). These patterns, as well as additional patterns disclosed herein, represent probable splicing variants which exist. The splicing variants are fully described in Goodearl et al., USSN 08/036,555, filed March 24, 1993, incorporated herein by reference.

More specifically, effects on cell communication may be achieved by contacting cells with a polypeptide defined by the formula

WYBAZCX

wherein WYBAZCX is composed of the polypeptide segments shown in Figure 13; wherein W comprises the polypeptide segment F, or is absent wherein Y comprises the polypeptide segment E, or is absent; wherein Z comprises the polypeptide segment G or is absent; and wherein X comprises the polypeptide segment C/D HKL, C/D H, C/D HL, C/D D, C/D' HL, C/D' HKL, C/D' HKL, C/D D' HKL, C/D D' HKL, C/D' D' HKL, C/D' D' HKL, C/D' D' HKL, C/D' D' HKL.

Furthermore, the invention includes a method of treating muscle cells by the application to the muscle cell of a

- -30kD polypeptide factor isolated from the MDA-MB 231 human breast cell line; or
- -35kD polypeptide factor isolated from the rat I-EJ transformed fibroblast cell line to the glial cell; or
 - -75kD polypeptide factor isolated from SKBR-3 human breast cell line; or
- -44kD polypeptide factor isolated from the rat I-EJ transformed fibroblast cell line; or
- -25kD polypeptide factor isolated from activated mouse peritoneal macrophages; or
 - -45kD polypeptide factor isolated from the MDA-MB 231 human breast cell; or
- -7 to 14kD polypeptide factor isolated from the ATL-2 human T-cell line to the glial cell; or
 - -25kD polypeptide factor isolated from the bovine kidney cells; or
 - -42kD ARIA polypeptide factor isolated from brain; or
 - -46-47kD polypeptide factor which stimulates 0-2A glial progenitor cells; or
- -43-45kD polypeptide factor, GGFIII, U.S. patent application Serial No. 07/931,041, filed August 17, 1992, incorporated herein by reference.

The invention includes use of any modifications or equivalents of the above polypeptide factors which do not exhibit a significantly reduced activity. For example, modifications in which amino acid content or sequence is altered without substantially adversely affecting activity are included. The statements of effect and use contained herein are therefore to be construed accordingly, with such uses and effects employing modified or equivalent factors being part of the invention.

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The human peptide sequences described above and presented in Figures 13, 14, 15, and 16, respectively, represent a series of splicing variants which can be isolated as full-length complementary DNAs (cDNAs) from natural sources (cDNA libraries

prepared from the appropriate tissues) or can be assembled as DNA constructs with individual exons (e.g., derived as separate exons) by someone skilled in the art.

Another aspect of the invention is the use of a pharmaceutical or veterinary formulation comprising any factor as defined above formulated for pharmaceutical or veterinary use, respectively, optionally together with an acceptable diluent, carrier or excipient and/or in unit dosage form. In using the factors of the invention, conventional pharmaceutical or veterinary practice may be employed to provide suitable formulations or compositions.

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A medicament is made by administering the polypeptide with a pharmaceutically effective carrier.

Thus, the formulations to be used as a part of the invention can be applied to parenteral administration, for example, intravenous, subcutaneous, intramuscular, intraorbital, ophthalmic, intraventricular, intracranial, intracapsular, intraspinal, intracisternal, intraperitoneal, topical, intranasal, aerosol, scarification, transdermal and by other slow release devices (i.e., osmotic pump-driven devices; see also U.S.S.N. 08/293,465, hereby incorporated by reference) and also oral, buccal, rectal or vaginal administration.

The formulations of this invention may also be administered by the transplantation into the patient of host cells expressing the DNA encoding polypeptides which are effective for the methods of the invention or by the use of surgical implants which release the formulations of the invention.

Parenteral formulations may be in the form of liquid solutions or suspensions; for oral administration, formulations may be in the form of tablets or capsules; and for intranasal formulations, in the form of powders, nasal drops, or aerosols.

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Methods well-known in the art for making formulations are to be found in, for example, "Remington's Pharmaceutical Sciences." Formulations for parenteral administration may, for example, contain as excipients sterile water or saline, polyalkylene glycols such as polyethylene glycol, oils of vegetable origin, or hydrogenated naphthalenes, biocompatible, biodegradable lactide polymer, or polyoxyethylene-polyoxypropylene copolymers may be used to control the release of the present factors. Other potentially useful parenteral delivery systems for the factors include ethylene-vinyl acetate copolymer particles, osmotic pumps, implantable

infusion systems, and liposomes. Formulations for inhalation may contain as excipients, for example, lactose, or may be aqueous solutions containing, for example, polyoxyethylene-9-lauryl ether, glycocholate and deoxycholate, or may be oily solutions for administration in the form of nasal drops, or as a gel to be applied intranasally. Formulations for parenteral administration may also include glycocholate for buccal administration, methoxysalicylate for rectal administration, or citric acid for vaginal administration.

The present factors can be used as the sole active agents, or can be used in combination with other active ingredients, e.g., other growth factors which could facilitate neuronal survival in neurological diseases, or peptidase or protease inhibitors.

The concentration of the present factors in the formulations of the invention will vary depending upon a number of issues, including the dosage to be administered, and the route of administration.

In general terms, the factors of this invention may be provided in an aqueous physiological buffer solution containing about 0.1 to 10% w/v compound for parenteral administration. General dose ranges are from about 1 µg/kg to about 1 g/kg of body weight per day; a preferred dose range is from about 0.01 mg/kg to 100 mg/kg of body weight per day. The preferred dosage to be administered is likely to depend upon the type and extent of progression of the pathophysiological condition being addressed, the overall health of the patient, the make up of the formulation, and the route of administration.

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A further general aspect of the invention is the use of a factor of the invention in the manufacture of a medicament, preferably for the treatment of a disease or disorder. The "GGF2" designation is used for all clones which were previously isolated with peptide sequence data derived from GGF-II protein (i.e., GGF2HBS5, GGF2BPP3) and, when present alone (i.e., GGF2 or rhGGF2), to indicate recombinant human protein encoded by plasmids isolated with peptide sequence data derived from the GGF-II protein (i.e., as produced in insect cells from the plasmid HBS5). Recombinant human GGF from the GGFHBS5 clone is called GGF2, rhGGF2 and GGF2HBS5 polypeptide.

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Methods for treatment of diseases or disorders using neuregulins in this manner are also part of the invention. Administration of neuregulins to induce the production of a substance or substances from a neuregulin responsive cell can be used in any disorder where an increase in a neuregulin inducible substance that is trophic for the disease

affected neurons would be of benefit. In peripheral nerve injury or peripheral nerve disorders such as the neuropathies administration of neuregulins will elicit the production of neurotrophic substances from known neuregulin target tissues such as Schwann cells and muscle. These induced substances can enhance axonal repair. Alzheimer's disease is another target for neuregulin therapy. In the brain, neuregulins are detectable in cholinergic motor neurons (Chen, et al., *J. Comparative Neurology* (1994) 349:389-400), these neurons degenerate in Alzheimer's disease and many show trophic responses to neurotrophic factors such as NGF. Neuregulins can be used to induce the synthesis of neurotrophic factors in those neurons that interact with cholinergic neurons. Similar therapeutic approaches may be used in other neurodegenerative disorders such as Parkinson's disease, amyotrophic lateral sclerosis, spinal muscular atrophy or any disease where stimulation of the synthesis of substances that are trophic for disease affected neurons might be of benefit.

Methods for treatment of diseases or disorders using nucleic acid constructs encoding neuregulins or neuregulin producer cells are also part of the invention.

Delivery of DNA to a cell or tissue that will take up the DNA, express the DNA and produce neuregulin as shown by Wolff et al., (Science (1990) 247:1465) and Ascadi et al., (Nature (1991) 352:815) is an aspect of the invention. The neuregulin produced by this method will act on the first cell type and elicit the responses described above. Genetic modification of cultured cells (or their precursors) such as fibroblasts (as shown by Wolff et al. Proc. Nat'l Acad. Sci. USA (1988) 86:1575) or such as those derived from the nervous system (as shown by Weiss et al. International Patent Application number PCT/US94/01053; publication number WO 94/16718) to induce the production of neuregulin from the cultured cells is another aspect of this invention. The genetically modified neuregulin producer cells can be transplanted to a position near the first cell type and elicit the responses described above.

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Assays for Determining Neuregulin Effect(s) on Cellular Communication

Described below are generic methods for detecting the ability of a neuregulin to induce in a first cell type, the production of a product (Product A) that is trophic for a second cell type. A general reference on cell and tissue culture is *Cell and Tissue Culture: Laboratory Procedures* (Ed. by A. Doyle, J. B. Griffiths, and D. G. Newell, John Wiley and Sons, New York, NY, 1994). General references on the culture of neural cells and tissues are *Methods in Neurosciences*, *Vol.* 2 (Ed. by P. M. Conn. Academic Press, Sand Diego, CA, 1990) and *Culturing Nerve Cells* (Ed. by G. Banker and K. Goslin, MIT Press, Cambridge, MA 1991). General references of immunocytochemistry are *Antibodies: A Laboratory Manual* (E. Harlow and D. Lane, Cold Spring Harbor Laboratory, Cold Spring Harbor, NY, 1988), and *Immunocytochemistry II* (Ed. by A. C. Cuello, John Wiley and Sons, New York, NY, 1993).

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The vertebrate cells used in this invention may be cultured in a variety of media. Commercially available media such as Ham's F10 (Sigma), Minimal Essential Medium ([MEM], Sigma), RPMI-1640 (Sigma), and Dulbecco's Modified Eagle's Medium ([DMEM], Sigma) are suitable for culturing the host cells. In addition, any of the media described in Ham and Wallace, Meth. Enz. (1979) 58:44; Barnes and Sato, Anal. Biochem. (1980) 102:255; U.S. Pat. Nos. 4,767,704; 4,657,866; 4,927,762; or 4,560,655; WO 90/03430; WO 87/00195 and U.S. Pat. Re. 30,985, may be used as culture media for the host cells. Any of these media may be supplemented as necessary with hormones and/or other growth factors (such as insulin, transferrin, or epidermal growth factor), salts (such as sodium chloride, calcium, magnesium, and phosphate), buffers (such as HEPES), nucleosides (such as adenosine and thymidine), antibiotics (such as Gentamycin™ drug), trace elements (defined as inorganic compounds usually present at final concentrations in the micromolar range), and glucose or an equivalent energy source. Any other necessary supplements may also be included at appropriate concentrations that would be known to those skilled in the art. The culture conditions, such as temperature, pH, and the like, are those previously used with the host cell selected for expression and will be apparent to the ordinarily skilled artisan.

Method I -

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The use of separate cultures of a first cell type and a second cell type, to demonstrate that neuregulin induces the first cell type to produce a secreted substance that is trophic for the second cell type.

- 1. Establish cultures of cells from the tissue of interest (e.g. spinal cord, pancreas, gut, etc.). These cultures are enriched for the first cell type such that preferably greater than 90% of the cells can be demonstrated to be the same cell type through the use of immunocytochemical and/or enzymatic markers (e.g. tubulin β3 for neurons (A. Banerjee, M. C. Roach, P. Trcka, and R. F. Luduena, Increased microtubule assembly in bovine brain tubulin lacking the type III isotype of b tubulin. *J. Biol. Chem.* (1990) 265:1794-1799), Islet-1 for pancreatic islet cells (O. Karlsson, S. Thor, T. Norbert, H. Ohlsson, and T. Edlund, Insulin gene enhanced binding protein Isl-1 is a member of a novel class of proteins containing both a homeo and a Cys-His domain. *Nature* (1990) 344:879-882)).
- 2. Establish cultures of cells from the same tissue of interest as in step 1. These cultures are enriched for the second cell type such that preferably greater than 90% of the cells can be demonstrated to be the same cell type through the use of immunocytochemical and/or enzymatic markers.
- 2. Expose the first cell type cultures to varying doses of neuregulin for varying periods of time, preferably greater than 1 minute and less than 7 days. At the end of the culture period, collect the conditioned culture medium, remove debris by centrifugation (200 g, 10 minutes) and filtration (nylon filter, 0.22 mm pore size). This medium (conditioned medium) will contain the secreted product(s) of the first cell type, Product A.
- 30 4. Replace or supplement the medium of the second cell type cultures with media prepared as in step 3. Include among these medium samples, medium that has been conditioned by the first cell type cultures in the absence of neuregulin (control conditioned medium). Include among the medium samples, media containing neuregulin that have not been conditioned by the first cell type cultures (non-conditioned medium).
 - 5. Maintain the second cell type cultures as described in step 4 for varying periods of time preferably greater than 1 day and less than 7 days. Assess various aspects of

cellular phenotype such as, but not limited to, cell survival, morphology, production of enzymes and secreted products, etc.

- 6. Assess the effects of the neuregulin. The neuregulin is trophic for the first cell type in a manner that promotes the production of products trophic for the second cell type if:
 - a. Medium conditioned by the first cell type cultures in the presence of neuregulin maintains or increases desired aspects of cellular phenotype such as, but not limited to cell survival, morphology, production of enzymes and secreted products, etc.;
- b. equal volumes of control conditioned medium lack the activity described in criterion (a.), or demonstrate lesser degrees of the activity described in criterion (a.);

and

c. equal volumes of non-conditioned medium lack the activity described in criterion (a.), or demonstrate lesser degrees of the activity described in criterion (a.).

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The induction by neuregulin of a secreted product, Product A, such that Product A affects a third cell type, can also be tested as in Method I. Establish cultures of cells from the same tissue of interest as in step 1. These cultures are enriched for the third cell type, such that preferably greater than 90% of the cells can be demonstrated to be the same cell type through the use of immunocytochemical and/or enzymatic markers. Substitute the third cell type cultures for the second cell type cultures in steps 4-6.

If Product A is not secreted, but is bound to the surface of the first cell type, or is bound to insoluble extracellular matrix associated with the first cell type, an alternative procedure is to be used:

Method II -

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The use of separate cultures of the first and second cell types, to demonstrate that neuregulin induces the first cell type to produce a substance on its surface that is trophic for the second cell type.

- 1. Establish cultures of cells from the tissue of interest (e.g. spinal cord, pancreas, gut, etc.). These cultures are enriched for the first cell type such that preferably greater than 90% of the cells can be demonstrated to be the same cell type through the use of immunocytochemical and/or enzymatic markers (e.g. tubulin β3 for neurons, Islet-1 for pancreatic islet cells).
- 2. Expose the first cell type cultures to varying doses of neuregulin for varying periods of time, preferably greater than 1 hour and less than 7 days. At the end of the culture period, remove the culture medium and establish a co-culture of the first and second cell types as follows. Rinse the first cell type cultures 3 times with fresh culture medium lacking neuregulin so as to rinse away residual neuregulin. Add back a suspension of the second cell type, from the same tissue of interest as in step 1 in fresh medium lacking neuregulin. The suspension is enriched for the second cell type, such that preferably greater than 90% of the cells can be demonstrated to be the same cell type through the use of immunocytochemical and/or enzymatic markers.
 - 3. In parallel to step 2, plate the same suspension of cells of the second cell type on the first cell type cultures that have not been treated with neuregulin (control co-cultures).
 - 4. Maintain the first cell type/second cell type co-cultures for varying periods of time preferably greater than 1 day and less than 7 days. Assess various aspects of cellular phenotype of the second cell type such as, but not limited to, cell survival, morphology, production of enzymes and secreted products, etc.
 - 5. Assess the effects of neuregulin. Neuregulin is trophic for the first cell type in a manner that promotes the production of products trophic for the second cell type if:
- a. The first cell type cultures pre-treated with neuregulin maintain or increase desired aspects of cellular phenotype of the second cell type such as, but not limited to cell survival, morphology, production of enzymes and secreted products, etc.;

and

b. The first cell type cultures that have not been pre-treated with Product A lack the activity described in criterion (a.), or demonstrate lesser degrees of the activity described in criterion (a.).

The induction by neuregulin of a cell surface-bound or extracellular matrix-bound product, Product A, such that Product A affects a third cell type, can also be tested as in Method II. In steps 2-4, use a suspension of the third cell type rather than the second cell type such that preferably greater than 90% of the cells can be demonstrated to be the third cell type through the use of immunocytochemical and/or enzymatic markers.

Described below are methods for detecting the activities of a neuregulin that induces neuronally-associated tissues to produce a neurotrophic product or product(s) (Product A):

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Method III -

The use of separate cultures of neurons and neuronally associated tissues, to demonstrate that neuregulin induces a neuronally associated tissue (the first cell type) to produce a secreted product that is trophic for neurons (the second cell type).

1. Establish neuron-free cultures of neuronally-associated cell types (e.g. glia, fibroblasts). These cultures are enriched for a single cell type (the first cell type) such that preferably greater than 90% of the cells can be demonstrated to be the same cell type through the use of immunocytochemical and/or enzymatic markers (e.g. S-100 for peripheral glia (K. R. Jessen and R. Mirsky, Schwann cell: early lineage, regulation of proliferation and control of myelin formation. *Curr. Op. Neurobiol.* (1992) 2:575-581), fibronectin for fibroblasts (K. M. Yamada, Cell surface interactions with extracellular materials. *Ann. Rev. Biochem.* (1983) 52:761-799)).

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- Establish cultures of neurons from the neuronal tissue of interest (e.g. superior cervical ganglion, spinal motor column). These cultures are enriched for neurons (the second cell type) such that preferably greater than 90% of the cells can be demonstrated to be the same cell type through the use of immunocytochemical and/or enzymatic markers (e.g. tubulin β3 for all neurons, choline acetyltransferase for cholinergic neurons (J.C. Martinou, A. L. V. Thai, G. Cassar, F. Roubinet, and M. J. Weber, Characterization of two factors enhancing choline acetyltransferase in cultures of purified rat motoneurons. J. Neurosci. (1989) 9:3645-3656)).
- 25 3. Expose the first cell type cultures to varying doses of neuregulin for varying periods of time, preferably greater than 1 hour and less than 7 days. At the end of the culture period, collect the conditioned culture medium, remove debris by centrifugation (200 g, 10 minutes) and filtration (nylon filter, 0.22 mm pore size).
- 30 4. Replace or supplement the medium of the second cell type cultures with media prepared as in step 3. Include among these medium samples, medium that has been conditioned by the first cell type cultures in the absence of neuregulin (control conditioned medium). Include among the medium samples, media containing neuregulin that have not been conditioned by the first cell type cultures (non-conditioned medium).
 - 5. Maintain the second cell type cultures as described in step 4 for varying periods of time preferably greater than 1 day and less than 7 days. Assess various aspects of

neuronal phenotype such as, but not limited to cell survival, neurite (axon or dendrite) outgrowth, neurotransmitter phenotype, etc.

- 6. Assess the effects of neuregulin. Neuregulin is trophic for neuronally-associated tissues in a manner that promotes the production of neurotrophic products if:
 - a. Medium conditioned by the first cell type cultures in the presence of neuregulin maintains or increases desired aspects of neuronal phenotype such as, but not limited to cell survival, increased neurite (axon or dendrite) outgrowth, neurotransmitter synthesis, etc.;
- b. equal volumes of control conditioned medium lack the activity described in criterion (a.), or demonstrate lesser degrees of the activity described in criterion (a.); and
 - c. equal volumes of non-conditioned medium lack the activity described in criterion (a.), or demonstrate lesser degrees of the activity described in criterion (a.).

If Product A is not secreted, but is bound to the surface of the first cell type, or is bound to insoluble extracellular matrix associated with the first cell type, an alternative procedure is to be used:

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Method IV -

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The use of separate cultures of neurons and neuronally associated tissues, to demonstrate that neuronally induces a neuronally associated tissue (the first cell type) to produce a substance on its surface that is trophic for neurons.

- 1. Establish neuron-free cultures of neuronally-associated cell types (e.g. glia, fibroblasts). These cultures are enriched for a single cell type (the first cell type) such that preferably greater than 90% of the cells can be demonstrated to be the same cell type through the use of immunocytochemical and/or enzymatic markers (e.g. S-100 for peripheral glia, fibronectin for fibroblasts).
- 2. Expose the first cell type cultures to varying doses of neuregulin for varying periods of time, preferably greater than 1 hour and less than 7 days. At the end of the culture period, remove the culture medium and establish a co-culture of the first cell type and neurons (the second cell type) as follows. Rinse the first cell type cultures 3 times with fresh culture medium lacking neuregulin so as to rinse away residual neuregulin. Add back a suspension of neurons from the neuronal tissue of interest (e.g. superior cervical ganglion, spinal motor column) in fresh medium lacking neuregulin. The suspension is enriched for the second cell type such that preferably greater than 90% of the cells can be demonstrated to be the same cell type through the use of immunocytochemical and/or enzymatic markers (e.g. tubulin β3 for all neurons, choline acetyltransferase for cholinergic neurons).
- 25 3. In parallel to step 2, plate the same suspension of the second cell type cells on the first cell type cultures that have not been treated with Product A (control co-cultures).
- 4. Maintain the first cell type/second cell type co-cultures for varying periods of time preferably greater than 1 day and less than 7 days. Assess various aspects of neuronal phenotype such as, but not limited to cell survival, neurite (axon or dendrite) outgrowth, neurotransmitter phenotype, etc.
- 5. Assess the effects of neuregulin. Neuregulin is trophic for the first cell type in a manner that promotes the production of products trophic for the second cell type if:
 - a. The first cell type cultures pre-treated with neuregulin maintain or increase desired aspects of neuronal phenotype such as, but not limited to cell survival, neurite (axon or dendrite) outgrowth, neurotransmitter phenotype, etc.;

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and

b. The first cell type cultures that have not been pre-treated with Product A lack the activity described in criterion (a.), or demonstrate lesser degrees of the activity described in criterion (a.).

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If cultures of non-neuronal cells of interest greater than 90% pure have not been established, the following method can be used:

Method V -

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The use of a mixed culture, to demonstrate that neuregulins induce the first cell type (neuronally associated cell types) to produce a product (Product A) that affects the second cell type.

- 1. Establish undissociated, explant cultures of the neuronal tissue of interest (e.g. superior cervical ganglion, spinal motor column). These cultures are not enriched for various cell types and are constituted of both neurons (the second cell type) and neuronally-associated cell types (the first cell type) as demonstrated through the use of immunocytochemical and/or enzymatic markers (e.g. tubulin \$3 for all neurons, acetylcholinesterase for cholinergic neurons, S-100 for peripheral glia, fibronectin for fibroblasts).
- 2. Expose explant cultures to varying doses of neuregulin for varying periods of time, preferably greater than 1 hour and less than 7 days. At the end of the culture period, assess various aspects of neuronal phenotype such as, but not limited to neuron survival, neurite (axon or dendrite) outgrowth, neurotransmitter phenotype, etc.
- 3. Establish cultures of neurons from the neuronal tissue of interest (e.g. superior cervical ganglion, spinal motor column). These cultures are enriched for neurons (the second cell type) such that preferably greater than 90% of the cells can be demonstrated to be the same cell type through the use of immunocytochemical and/or enzymatic markers (e.g. tubulin β3 for all neurons, choline acetyltransferase for cholinergic neurons).
 - 4. Expose the second cell type cultures to varying doses of neuregulin for varying periods of time, preferably greater than 1 hour and less than 7 days. At the end of the culture period, assess various aspects of neuronal phenotype such as, but not limited to neuron survival, neurite (axon or dendrite) outgrowth, neurotransmitter phenotype, etc.
 - 5. Assess the effects of neuregulin. Neuregulin is trophic for neuronally-associated tissues in a manner that promotes the production of neurotrophic products if:
- a. in explant cultures the presence of neuregulin maintains or increases desired aspects of neuronal phenotype such as, but not limited to neuron survival, neurite (axon or dendrite) outgrowth, neurotransmitter synthesis, etc.;

and

b. in the second cell type cultures, neuregulin lacks the activity described in criterion (a.), or demonstrates lesser degrees of the activity described in criterion (a.)

EXAMPLES

Example 1

The Effect of Recombinant Human Glial Growth Factor 2 on Sympathetic Ganglion Outgrowth in an *In Vitro* Model of Peripheral Nerve Gap Entubulation

Purpose

One approach to the repair of injuries in which a peripheral nerve has been severed is to suture the nerve endings together via a biocompatible tube, a procedure referred to as entubulation. The tube may be filled with various agents thought to improve the growth and regeneration of the nerve. Peripheral nerves contain a variety of cell types: neurons (or more appropriately, the axons emanating from neuron cell bodies located in the spinal cord and associated ganglia), Schwann cells (peripheral glia), fibroblasts, and resident macrophages. Axons regenerate from the side of the nerve gap proximal to the spinal cord and associated ganglia; other cell types contribute to regeneration by migrating in from both sides of the gap and proliferating.

In an effort to devise an *in vitro* model of entubulation, a technique was developed in which fragments of the rat superior cervical ganglion (SCG) are cultured in segments of surgical tubing used in whole animal models of peripheral nerve entubulation. SCG neurons are homogenous in their trophic requirements and project axons exclusively through peripheral nerves; SCG fragments also contain Schwann cells, fibroblasts, and macrophages. In this model the SCG fragments serve as surrogate proximal nerve endings, and the outgrowth of axons and supporting cell types can be observed in a simplified environment. The focus of this example was to examine the effects of rhGGF2 on Schwann cell and axon behavior in this *in vitro* model of peripheral nerve entubulation.

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Methods and Materials

Tube Preparation

Tubing used for this study was polyethylene tubing with an internal diameter of 1.19 mm and outer diameter of 1.70 mm (Intramedic®: Becton Dickinson and Company; Parsippany, New Jersey). A length of tubing somewhat longer than actually needed was cut in a sterile tissue culture hood, immersed in 70% ethanol, and flushed repeatedly with 70% ethanol using a syringe with a 19-gauge needle. After soaking the

tubing for approximately 30 minutes, it was flushed again with air, and allowed to dry in the hood. After drying, the tubing was cut into 10 mm segments with a sterile scalpel, and stored in a sterile Petri dish.

5 Culture Medium

Culture medium was made freshly on the day of culture assembly. All components were kept cold (either 4° C or on ice), as was the final solution until culture assembly was completed.

10	Sterile Water	2.60					
	Sodium bicarbonate (2% w/v)	1.50					
	Penicillin/Streptomycin stock*	0.15					
	L-Glutamine (200 mM)	0.15					
15	Fetal Bovine Serum**	0.75					
	Sodium hydroxide (0.1 M)	0.90					
	10x Medium***	1.50					
	Collagen solution****	<u>7.40</u>					
	TOTAL	15.00					

TOTAL 15.00 ml

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Medium was used as is, or was supplemented with rhGGF2 as indicated.

^{* 5000} units/ml penicillin, 5 mg/ml streptomycin

^{**} heat inactivated (Hyclone; Logan, UT)

^{***} one packet of low glucose Dulbecco's Modified Essential Medium (DMEM: Gibco/BRL; Grand Island, NY) meant to make 1 liter of medium dissolved in 100 ml of sterile water

^{**** 3} mg/ml Vitrogen-100® (Celtrix Pharmaceuticals; Santa Clara, CA)

Tube Culture Assembly

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A schematic diagram of culture assembly is shown in Figure 1. SCGs were dissected from postnatal day 0-2 rats on the day of assembly, cleaned of connective tissues and proximal nerve stumps, bisected, and stored in physiological saline at 4° till needed. Since the collagen-containing medium gels at room temperature or higher, it is necessary to assemble the cultures in 4° cold room. Working with watchmaker's forceps under a dissection microscope at total magnification of 8x, individual segments of cleaned tubing were picked up and filled with culture medium using a syringe with a 27-gauge needle. A single piece of bisected SCG was then placed at the very end of each tube, and each tube placed in an individual well of a 24-well tissue culture plate. Only the central eight wells of a 24-well plate were used for tube placement, and the remaining wells were filled with sterile water to maintain plate humidity. The plate was then placed in a 37°, 10% carbon dioxide incubator. After allowing the cultures to gel and equilibrate with the incubator atmosphere, the plates were sealed with paraffin film to further protect the culture assemblies from dehydration, and returned to the incubator until preparation for immunocytochemistry and analysis.

Immunocytochemistry

After 2, 5, and 10 days in vitro, the contents of individual tube cultures were extruded into phosphate buffered saline (PBS) using a PBS-filled syringe with a bluntended 18-gauge needle. The collagen gels retained structural integrity and were fixed in 4% paraformaldehyde in PBS for 30 minutes at room temperature. After 3 washes with PBS, the cultures were blocked in 1% goat serum/0.1% Triton X-100 in PBS for 30 minutes. After blocking, the solution was changed to 1% goat serum in PBS (GPBS) containing 1:4 rabbit anti-S-100 (a Schwann cell marker; Incstar; Stillwater, MN) and 1:400 mouse anti-tubulin 83 (an axon marker; Sigma; St. Louis, MO). After incubating the samples in primary antibody for 1 hour at room temperature, they were washed 3 times with PBS, and incubated for an additional hour in GPBS containing 1:200 peroxidase-conjugated goat anti-mouse immunoglobulin, and 1:200 alkaline phosphatase-conjugated goat anti-rabbit immunoglobulin (Pierce Chemical; Rockford, IL). The samples were then washed 3 times with PBS, and the stains developed. The S-100 was first developed using stable pre-mixed NBT/BCIP (Gibco/BRL) to yield a blue stain, and after rinsing with PBS, the tubulin $\beta 3$ was developed using 3-amino-9ethylcarbazole (AEC; Sigma) per manufacturer's directions. After final rinsing with PBS, the samples were mounted on microscope slides using aqueous mounting medium. After the mounting medium dried, the cellular outgrowth could be analyzed.

Scoring and Analysis

As schematized in Figure 2, a grid reticule was placed in the microscope ocular, and at a total magnification of 160x, the total number of S-100+ Schwann cells in each column (referred to as "bins") was counted. Each bin has a width of 50 mm as determined using a stage micrometer. And as noted, the number of tubulin $\beta3^+$ neurites intersecting every vertical line was also counted. The actual grid was not large enough to cover the entire length of cellular outgrowth and was shifted along as needed by translational movement of the microscope stage. All data points represent the average \pm the standard error of the mean (n = 6 to 7 for every data point).

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Results

First shown is the analysis of Schwann cell number as a function of distance from the SCG explant (Figure 3A-D). It is clear that the presence of rhGGF2 affects the behavior of Schwann cells relative to the control condition. There does not appear to be any difference among the 3 doses of rhGGF2. Generally, by 5 days in rhGGF2, there is a large increase in the number of Schwann cells proximal to the explant, but the Schwann cells appear to have moved only about as far as they have in the control case (somewhat further at the highest dose). By 10 days in rhGGF2 the overall number of Schwann cells has decreased, but the cells still present have definitely migrated farther than in the absence of rhGGF2. In the absence of rhGGF2, the controls look no different between days 5 and 10. The total number of Schwann cells in the various conditions is shown in Figure 4. Again, there is a decrease in cell number at day 10, but there is no obvious difference between the different doses of rhGGF2. The day 10 tubes contain more debris, and this is probably due to cell death. This is due to the culture situation since 10 days appears to be the longest that one can maintain these tube cultures without overt signs of dehydration and nutrient depletion in the limited volume of culture medium (approximately 10 µl per tube).

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A difference is apparent when neurites are scored in the various doses of rhGGF2 (Figure 5). At doses of rhGGF2 greater than or equal to 50 ng/ml, a profound increase takes place in the number of neurites and the extent to which they have grown away from the explant.

Discussion and Conclusions

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This study demonstrates that the dose range in which there are observable effects on Schwann cell proliferation and emigration from the explant is different from that which causes a major increase in neurite outgrowth. In the case of the former, it

appears that the effect has plateaued at the lowest dose tested, 5 ng/ml. As for the requirement of ≥50 ng/ml rhGGF2 to boost neurite regeneration, there are two possible mechanisms to account for this. One is that rhGGF2 is acting directly upon the neurons, and the other is that rhGGF2 induces a non-neuronal cell type to produce a neurite promoting factor (e.g. NGF, secreted extracellular matrix proteins, proteases, and/or protease inhibitors). As is demonstrated in Example 2, the first hypothesis is not likely since rhGGF2 has no effect upon neuronal survival or outgrowth in low density cultures of dissociated SCG neurons. This lack of a direct effect on neurons implies that the rhGGF2 promotion of neurite outgrowth is due to rhGGF2 induced production of neurite promoting factors by non-neuronal cells.

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Example 2

The Promotion of Axon Outgrowth by Recombinant Human Glial Growth Factor 2 is Not Due to a Direct Effect on Neurons

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Purpose

As demonstrated in Example 1, rhGGF2 not only promotes Schwann cell proliferation and migration in an *in vitro* model of peripheral nerve entubulation, but also promotes robust axonal outgrowth. To test whether this may be due to direct effects of rhGGF2 on SCG neurons, low density cultures of dissociated SCG neurons were established in which the effects of rhGGF2 could be examined. SCG neurons are normally dependent upon nerve growth factor (NGF) for survival, so rhGGF2 was tested for direct neuronal effects in the simultaneous presence of a wide range of NGF concentrations.

Methods and Materials

Cell Culture

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SCGs were dissected from postnatal day 0-2 rats, cleaned of connective tissue and proximal nerve stumps, and dissociated by enzymatic digestion and trituration. Enzymatic digestion was performed using 1 mg/ml trypsin (Sigma; St. Louis, MO) and 1 mg/ml collagenase (Boehringer-Mannheim; Indianapolis, IN) in calcium- and magnesium-free Hanks's Balanced Salt Solution (HBSS; Gibco/BRL; Grand Island, NY), for 1 hour at 37° C. Trituration was performed using a flame-polished Pasteur pipet. Dissociated neurons were taken up in plating medium and pre-plated in tissue culture dishes for 1 hour to remove the majority of the rapidly adherent, non-neuronal cells. Plating medium consisted of low glucose DMEM (Gibco/BRL) supplemented with glutamine, penicillin/streptomycin, and fetal bovine serum to the same concentrations as described in Example 1. Non-adherent cells (primarily neurons), were pelleted by centrifugation and resuspended in plating medium. These cells were finally plated at a density of 5000 cells per well in collagen-coated, 24-well plates such that the cells were exposed to a 2-dimensional dose-response matrix of NGF and rhGGF2 (Figure 6A). Plates were set up in duplicate on 2 different dates; at the completion of both experiments N = 4 for each of the 24 conditions. The cultures were only allowed to progress for 2 days since this is a time frame in which any

contaminating Schwann cells could have only undergone a single doubling, and sufficient for ascertaining whether the factors have promoted neuronal survival.

Staining and Scoring of the Cultures

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After 2 days, the cultures were fixed and stained for tubulin \$3 as described in Example 1. The tubulin \$3-positive, neurite-bearing cells were counted in each well at a total magnification of 100x. Due to meniscus effects, and incubator vibration during the initial plating period, cells tend to preferentially concentrate in the center of the well. Thus in order to get a reasonably representative count of cell number, 5 fields per well were counted: the center most field and four flanking fields (Figure 6B). This manner of counting was used on all wells and is sufficiently consistent for the purpose of comparing the effects of different growth factor concentrations and combinations. The number of cells counted in every well was normalized such that the average number counted in the wells that received 0 ng/ml rhGGF2, and 100 ng/ml NGF equals a value of 100.

Results

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It is clear from the results presented in Figure 7, that rhGGF2 has no direct effect on the survival of SCG neurons. All surviving neurons exhibited robust axon outgrowth, and there was no noticeable effect on the extent of axon outgrowth. As expected in the absence of rhGGF2, the number of neurons reaches a plateau at 10 ng/ml. The presence or absence of rhGGF2 appears to make no difference at the 3 doses tested.

Discussion and Conclusions

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In light of the results presented in Example 1, it was necessary to examine whether the effect of rhGGF2 on axon outgrowth could be attributed to a direct effect of rhGGF2 on the neurons in question. The results of Example 2 make it clear that this is not the case. Thus one must conclude that the effect of rhGGF2 on axon outgrowth observed in the tube paradigm is due to a "bystander effect" rather than a direct action on the neurons. Thus rhGGF2 can promote the healing response of injured neurons by inducing the production of neurite promoting factors by non-neuronal support cells.

Example 3

Increase in myelinated axon growth in an animal model of peripheral nerve injury mediated by a neuregulin

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An animal model of peripheral nerve repair was used to test the ability of a neuregulin (rhGGF2) to increase the number of regenerating axons. The rationale is that added rhGGF2 will induce increases in Schwann cell (the first cell type) numbers as well as increases in the levels of trophic factors (Product A) produced by Schwann cells that, in turn, will affect a second cell type, the regenerating axons (the second cell type) as measured by increases in the number of myelinated axons (response).

Fisher 344 rats (male, 195-250g) were surgically prepared and one sciatic nerve was transected resulting in a 10mm gap. Polyethylene guide tubes (13mm in length, 1.1mm internal diameter) were prepared. These tubes contained a flat sliver of a collagen coated Immobilon filter (1.0x10mm) containing immobilized rhGGF2 and were prepared as described in U.S. Patent Application Serial No. 08/293,465, filed on August 19, 1994, hereby incorporated by reference (Immobilon: Millipore, Corp., Bedford, MA). The strips were inserted into the lumen of the guide tubes. rhGGF2 was used at a concentration of 162 μ g/ μ L (in phosphate buffered saline), 2.5 μ L of this solution was added per strip. Control tubes were prepared containing collagen coated Immobilon strips treated with phosphate buffered saline alone. Tubes were secured with a single suture at the proximal and distal ends after filling the lumen with physiological saline and sealing the ends with vaseline.

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Animals (10 rhGGF2 treated, 10 controls) were sacrificed at 28 days and the section of sciatic nerve containing the tube was excised, the nerve was removed from the tube and a cross section was taken from the mid point of the tube and prepared for histological analysis. The material was fixed in 4% paraformaldehyde and 2% glutaraldehyde for 24h and then post fixed in 2% osmium tetroxide and embedded in glycomethacrylate. One micron cross sections were taken and stained with $1\mu M$ toluidine blue.

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A histological analysis of a section from the mid point of the tube was performed and measurements were made of the total number of myelinated axons in a section and the total endoneurial area in each section. The data are shown in Figures 8A and 8B.

The rhGGF2 treated animals showed a 2.1 fold increase in the number of myelinated axons over the control animals.

The results of this study demonstrate a positive effect of exogenously added rhGGF2 on the growth of myelinated axons. In consideration of the data discussed in Example 1 where rhGGF2 acts on Schwann cells to induce the synthesis of products that are trophic for regenerating axons in an *in vitro* paradigm it is concluded that a similar mechanism is responsible for the rhGGF2 mediated enhancement of the growth of axons *in vivo*.

Claims:

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- 1. A method of affecting cellular communication in a vertebrate, comprising administration of a neuregulin to said vertebrate wherein said neuregulin interacts with a first cell type, resulting in production of a Product A of said first cell type and said Product A affects a function of a second cell type.
- A method of Claim 1, wherein, said affect in function of said second cell type results
 in the production of a Product B of said second cell type and said Product B affects the function of said first cell type or a third cell type.
 - 3. A method of affecting cellular communication in a vertebrate, comprising administration of a neuregulin-producing cell to said vertebrate wherein a neuregulin is produced and said neuregulin interacts with a first cell type, resulting in production of a Product A of said first cell type and said Product A affects a function of a second cell type.
- 4. A method of Claim 3, wherein, said affect in function of said second cell type results
 in the production of a Product B of said second cell type and said Product B affects the function of said first cell type or a third cell type.
 - 5. A method of affecting cellular communication in a vertebrate, comprising administration of DNA encoding a neuregulin to said vertebrate wherein DNA is incorporated into a genome of a cell and said DNA is expressed in said cell resulting in the production of said neuregulin which interacts with a first cell type, resulting in production of a Product A of said first cell type and said Product A affects a function of a second cell type.
- 6. A method of Claim 5, wherein, said affect in function of said second cell type results in the production of a Product B of said second cell type and said Product B affects the function of said first cell type or a third cell type.
 - 7. A method of Claim 1 wherein said vertebrate is a human.
 - 8. A method of Claim 1 wherein said first cell type is a nervous system support cell.
 - 9. A method of Claim 8 wherein said nervous system support cell is a Schwann cell.

- 10. A method of Claim 1 wherein said first cell type is a neuron.
- 11. A method of Claim 1 wherein said first cell type is a muscle cell.

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- 12. A method of Claim 1 wherein said Product A is a neurotrophic agent.
- 13. A method of Claim 1 wherein said Product A is a matrix molecule.
- 10 14. A method of Claim 1 wherein said Product A is a protease.
 - · 15. A method of Claim 1 wherein said Product A is a protease inhibitor.
 - 16. A method of Claim 1 wherein said second cell type is a nervous system cell.

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- 17. A method of Claim 1 wherein said second cell type is a muscle cell.
- 18. A method of Claim 1 wherein said affect in function of said second cell type is differentiation.

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- 19. A method of Claim 1 wherein said affect in function of said second cell type is mitosis.
- 20. A method of Claim 1 wherein said affect in function of said second cell type is survival.
 - 21. A method of Claim 2 wherein said Product B is a neurotrophic agent.
 - 22. A method of Claim 2 wherein said Product B is a matrix molecule.

- 23. A method of Claim 2 wherein said Product B is a protease.
- 24. A method of Claim 2 wherein said Product B is a protease inhibitor
- 35 25. A method of Claim 2 wherein said Product B is a neuregulin.
 - 26. A method of Claim 25 wherein said neuregulin is rhGGF2.

27. A method of Claim 2 wherein said affect in function of said first cell type or said third cell type is differentiation.

- 28. A method of Claim 2 wherein said affect in function of said first cell type or said third cell type is mitosis.
 - 29. A method of Claim 2 wherein said affect in function of said first cell type or said third cell type is survival.
- 10 30. A method of Claim 2 wherein said third cell type is a nervous system cell.
 - 31. A method of Claim 2 wherein said third cell type is a muscle cell.
- 32. A method of treating a neurological disorder in a mammal, comprising administration of a therapeutically effective amount of a neuregulin to said mammal wherein said neuregulin interacts with a nervous system cell, resulting in the production of a neurotrophic agent which affects the function of a neuron cell type.
- 33. A method of treating a neurological disorder in a mammal, comprising administration of a neurogulin producing cell to said mammal wherein said produced neurogulin interacts with a nervous system cell, resulting in the production of a neurotrophic product which affects the function of a neuron cell type.
- 34. A method of treating a neurological disorder in a mammal, comprising administration of DNA encoding a neurogulin to said mammal wherein said neurogulin is produced and interacts with a nervous system cell, resulting in the production of a neurotrophic agent which affects the function of a neuron cell type.
- 35. A method of treating peripheral neuropathy, amyotrophic lateral sclerosis, spinal muscular atrophy, nerve injury, Alzheimer's Disease, Parkinson's Disease and spinal cord injury comprising the administration of a therapeutically effective amount of a neuregulin wherein said neuregulin interacts with a first cell type, resulting in the production of a Product A of the first cell type and said Product A affects a function of a second cell type.
 - 36. A method of inducing the endogenous production of a product by a cell in a vertebrate comprising administration of a neuregulin to said vertebrate, wherein said cell produces said product.

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37. A method of claim 36 wherein said product is a neurotrophic agent.

Figure 1

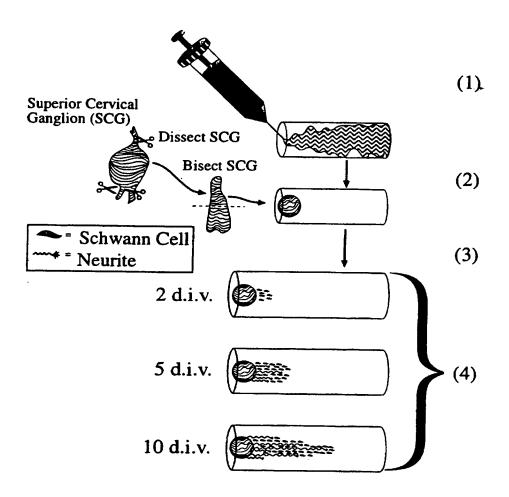


Figure 2

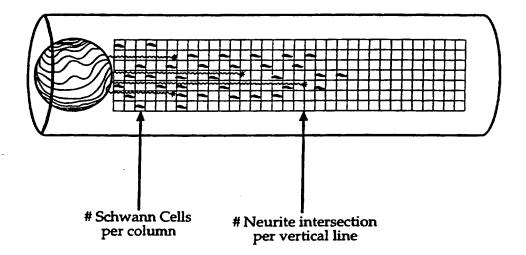
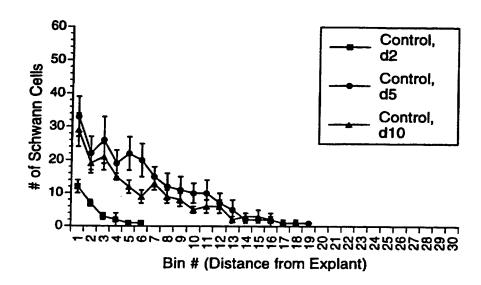


Figure 3A



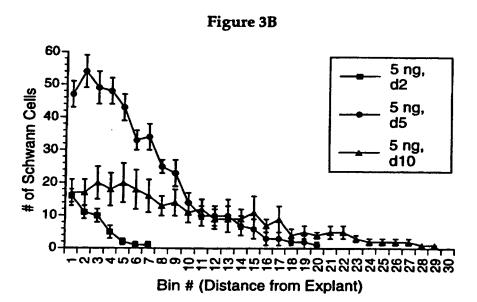


Figure 3C

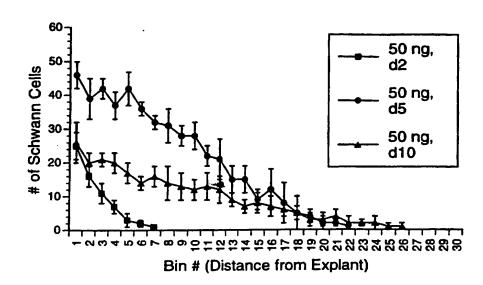


Figure 3D

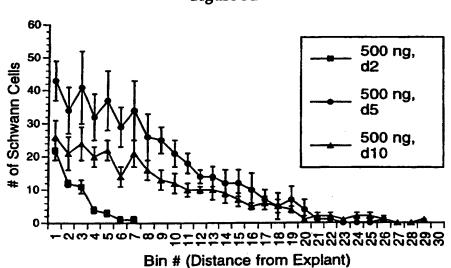
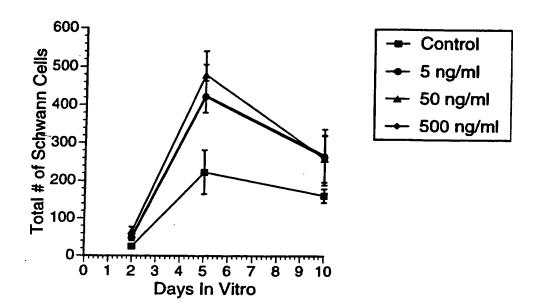


Figure 4



WO 96/15812

Figure 5

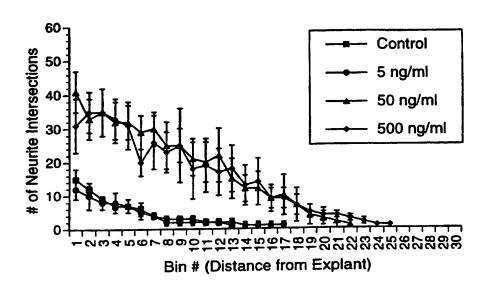


Figure 6A

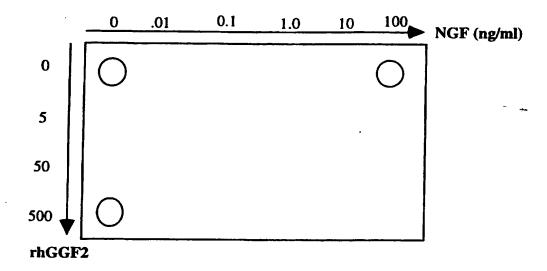


Figure 6B

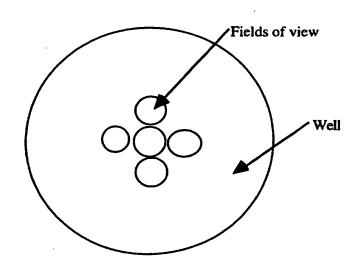


Figure 7

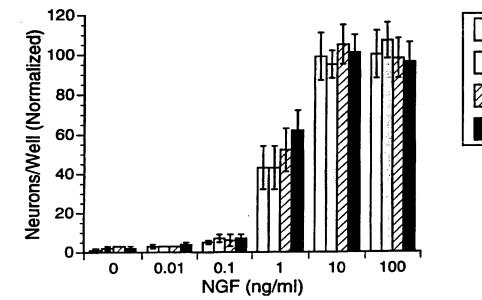




Figure 8A

	Total Myelinated Axons	Total Endoneurial Area	Axon Density
GGF group (n=9)			
GGF2 mean	741	165345(μM²)	4143(per mm ²)
GGF2 std. error	200	14809	951
Control group (n=10)			
Control mean	355	130931(μM²)	2938(per mm²)
Control std. error	76	14917	722

Figure 8B

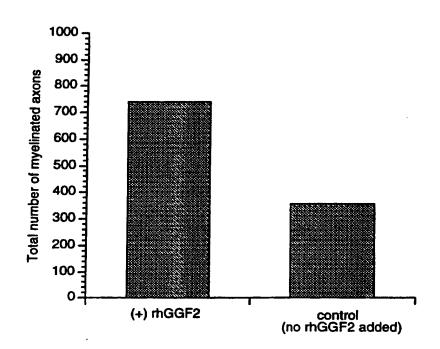
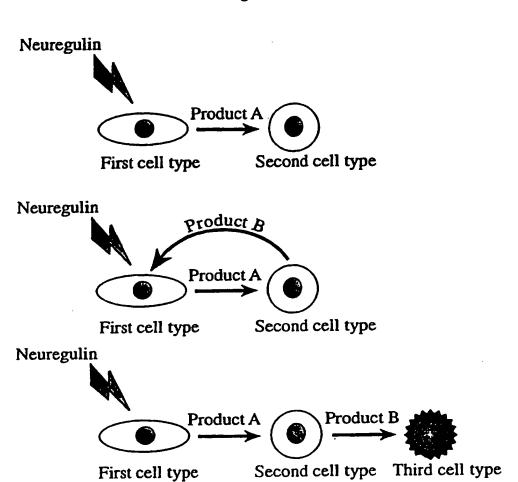
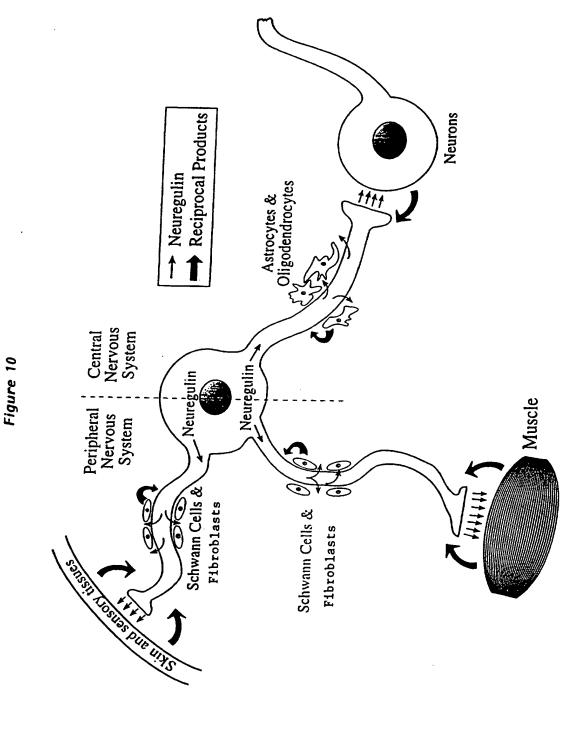


Figure 9





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Figure 11 A

CCTG	CAG	CAT His 1	CAA Gln	GTG Val	TGG Trp	GCG Ala 5	GCG Ala	AAA Lys	GCC Ala	GGG Gly	GGC Gly 10	TTG Leu	AAG Lys	AAG Lys	GAC Asp	TCG Ser 15	CTG Leu	55
CTC Leu	ACC Thr	GTG Val	CGC Arg 20	CTG Leu	GGC Gly	GCC Ala	TGG Trp	GGC Gly 25	CAC His	CCC Pro	GCC Ala	TTC Phe	CCC Pro 30	TCC Ser	TGC Cys			103
GGG Gly	CGC Arg	CTC Leu 35	AAG Lys	GAG Glu	GAC Asp	AGC Ser	AGG Arg 40	TAC Tyr	ATC Ile	TTC Phe	TTC Phe	ATG Met 45	GAG Glu	CCC Pro	GAG Glu			151
GCC Ala	AAC Asn 50	AGC	AGC Ser	GGC Gly	GGG Gly	CCC Pro 55	GGC	CGC Arg	CTT Leu	CCG Pro	AGC Ser 60	CTC Leu	CTT Leu	CCC Pro	CCC Pro			199
TCT Ser 65	CGA Arg	GAC Asp	GGG Gly	CCG Pro	GAA Glu 70	CCT Pro	CAA Gln	GAA Glu	GGA Gly	GGT Gly 75	CAG Gln	CCG Pro	GGT Gly	GCT Ala	GTG Val 80			247
CAA Gln.	CGG Arg	TGC Cys	GCC Ala	TTG Leu 85	CCT Pro	CCC Pro	CGC Arg	TTG Leu	AAA Lys 90	GAG Glu	ATG Met	AAG Lys	AGT Ser	CAG Gln 95	GAG Glu			295
TCT Ser	GTG Val	GCA Ala	GGT Gly 100	TCC Ser	AAA Lys	CTA Leu	Val	CTT Leu 105	CGG Arg	TGC Cys	GAG Glu	Thr	AGT Ser L10	TCT Ser	GAA Glu			343
TAC Tyr	TCC Ser	TCT Ser 115	CTC Leu	AAG Lys	TTC Phe	AAG Lys	TGG Trp 120	TTC Phe	AAG Lys	AAT Asn	GGG Gly	AGT Ser 125	GAA Glu	TTA Leu	AGC Ser			391
CGA Arg	AAG Lys 130	AAC Asn	AAA Lys	CCA Pro	GAA Glu	AAC Asn 135	ATC Ile	AAG Lys	ATA Ile	CAG Gln	AAA Lys 140	AGG Arg	CCG Pro	GGG Gly	AAG Lys			439
TCA Ser 145	GAA Glu	CTT Leu	CGC Arg	ATT Ile	AGC Ser 150	AAA Lys	GCG Ala	TCA Ser	CTG Leu	GCT Ala 155	GAT Asp	TCT Ser	GGA Gly	GAA Glu	TAT Tyr 160			487
ATG Met	TGC Cys	AAA Lys	GTG Val	ATC Ile 165	AGC Ser	AAA Lys	CTA Leu	GGA Gly	AAT Asn 170	GAC Asp	AGT Ser	GCC Ala	TCT Ser	GCC Ala 175	AAC Asn			535
ATC Ile	Thr	Ile	GTG Val 180	Glu	Ser	Asn	Gly	Lys	Arg	Cys	Leu	Leu	CGT Arg 190	GCT Ala	ATT			583
TCT Ser	CAG Gln	TCT Ser 195	CTA Leu	AGA Arg	GGA Gly	GTG Val	ATC Ile 200	AAG Lys	GTA Val	TGT Cys	GGT Gly	CAC His 205	ACT Thr					625
TGAA	TCAC	GC A	AGGT	STGTC	A A	ATCTO	TATI	TG	AACA	ATA	AAA	ATCA'	TGA	AAGG	AAAA	AA		685
AAAA	XAAA	LAA A	AATCO	ATGI	C GZ	CTC	AGAT	r GTY	GCT	GCAG	GTC	GACT	CTA	GAGG	ATCC	С		744

Figure 11 B

CC	IGCA	Hi 1	s Gl	n Va	G TG	g GCC p Ala 5	G GCC	Lys	A GCC	C GG(a Gly	G GGC 7 Gly 10	TTO Let	G AAG 1 Lys	S AAC S Lys	GAC Asp	TCG Ser 15	CTG Leu	55
CT Le	C AC	C GI r Va	G CG 1 Ar 20	C CTO	G GGG	C GCC V Ala	Trp	GGC Gly 25	CAC His	C CCC	GCC Ala	TTC Phe	C CCC Pro 30	TCC Ser	TGC Cys			103
GG(G CGG	C CT g Le 35	u ry:	G GAO s Glu	G GAC	C AGC Ser	AGG Arg 40	TAC	ATC Ile	TTC Phe	TTC Phe	ATG Met	GAG Glu	CCC Pro	GAG Glu			151
GC0 Ala	2 AAC 2 Lys 50	C AG	C AGC	GGC Gly	GGG Gly	CCC Pro 55	GGC Gly	CGC Arg	CTI Leu	CCG Pro	AGC Ser 60	CTC Leu	CTT Leu	CCC	CCC Pro			199
TCI Ser 65	CGA Arg	GA(Asp	GGG Gly	CCG Pro	GAA Glu 70	CCT Pro	CAA Gln	GAA Glu	GGA Gly	GGT Gly 75	CAG Gln	CCG Pro	GGT Gly	GCT Ala	GTG Val 80			247
CAA Gln	. CGG . Arg	TGC Cys	GCC Ala	TTG Leu 85	CCT Pro	CCC Pro	CGC Arg	TTG Leu	AAA Lys 90	GAG Glu	ATG Met	AAG Lys	AGT Ser	CAG Gln 95	GAG Glu			295
TCT Ser	GTG Val	GCA Ala	GGT Gly 100	TCC Ser	AAA Lys	CTA Leu	Val	CTT Leu 05	CGG Arg	TGC Cys	GAG Glu	Thr	AGT Ser L10	TCT Ser	GAA Glu			343
TAC Tyr	TCC Ser	TCT Ser 115	CTC Leu	AAG Lys	TTC Phe	AAG Lys	TGG Trp 120	TTC Phe	AAG Lys	AAT Asn	Gly	AGT Ser 125	GAA Glu	TTA Leu	AGC Ser			391
CGA Arg	AAG Lys 130	AAC Asn	AAA Lys	CCA Gly	GAA Gly	AAC Asn 135	ATC . Ile :	AAG Lys	ATA Ile	Gln	AAA Lys 140	AGG Arg	CCG Pro	GGG Gly	AAG Lys			439
TCA Ser 145	GAA Glu	CTT Leu	CGC Arg	ATT Ile	AGC Ser 150	AAA Lys .	GCG :	ICA (Ser :	Leu	GCT Ala 155	GAT (TCT Ser	GGA Gly	Glu	TAT Tyr 160			487
ATG Met	TGC Cys	AAA Lys	val	ATC . Ile . 165	AGC Ser	AAA (Lys)	CTA (Leu (Sly /	AAT Asn 170	GAC ASD	AGT (Ser)	GCC Ala	Ser .	GCC Ala	AAC Asn			535

Figure 11 B'

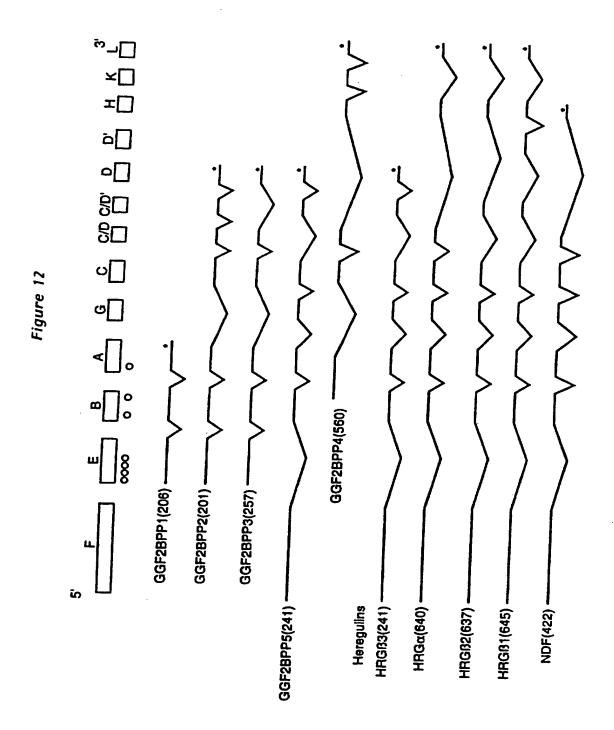
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AGC CAT CTT GTC AAG TGT GCA GAG AAG GAG AAA ACT TTC TGT GTG AAT Ser His Leu Val Lys Ser Ala Glu Lys Glu Lys Thr Phe Cys Val Asn 195 200 205	631
GGA GGC GAG TGC TTC ATG GTG AAA GAC CTT TCA AAT CCC TCA AGA TAC Gly Glu Cys Phe Met Val Lys Asp Leu Ser Asn Pro Ser Arg Tyr 210 215	679
TTG TGC AAG TGC CAA CCT GGA TTC ACT GGA GCG AGA TGT ACT GAG AAT Leu Cys Lys Cys Gln Pro Gly Phe Thr Gly Ala Arg Cys Thr Glu Asn 235	727 _.
GTG CCC ATG AAA GTC CAA ACC CAA GAA AGT GCC CAA ATG AGT TTA CTG Val Pro Met Lys Val Gln Thr Gln Glu Ser Ala Gln Met Ser Leu Leu 255	775
GTG ATC GCT GCC AAA ACT ACG TAATGGCCAG CTTCTACAGT ACGTCCACTC Val Ile Ala Ala Lys Thr Thr 260	826
CCTTTCTGTC TCTGCCTGAA TAGCGCATCT CAGTCGGTGC CGCTTTCTTG TTGCCGCATC	886
TCCCCTCAGA TTCCTCCTAG AGCTAGATGC GTTTTACCAG GTCTAACATT GACTGCCTCT	946
GCCTGTCGCA TGAGAACATT AACACAAGCG ATTGTATGAC TTCCTCTGTC CGTGACTAGT	1006
GGGCTCTGAG CTACTCGTAG GTGCGTAAGG CTCCAGTGTT TCTGAAATTG ATCTTGAATT	1066
ACTGTGATAC GACATGATAG TCCCTCTCAC CCAGTGCAAT GACAATAAAG GCCTTGAAAA	1126
GTCAAAAAA AAAAAAAAAA AAAAAATCGA TGTCGACTCG AGATGTGGCT GCAGGTCGAC	1193
TCTAGAG	

Figure 11 C

CC	rgca(G CA' Hi: 1	T CA	A GT n Va	G TG(l Tr _l	G GCG Ala 5	G GCG A Ala	AAA Lys	A GCC S Ala	C GGC a Gly	GGC Gly 10	Leu	AAG Lys	AAG Lys	GAC Asp	TCG Ser 15	CTG Leu	55
CT	C ACC	C GTY	G CGC l Arg 20	C CT	G GG(GCC Ala	TGG Trp	GGC Gly 25	CAC His	C CCC	GCC	TTC Phe	CCC Pro	TCC Ser	TGC Cys			103
GG(G CGC / Arg	CTC Let 35	AAG Lys	G GAG	GAC Asp	AGC Ser	AGG Arg 40	TAC Tyr	ATC	TTC Phe	TTC Phe	ATG Met 45	GAG Glu	CCC Pro	GAG Glu			151
GCC Ala	AAC Asn 50	AGC Ser	AGC Ser	GGC Gly	GGG Gly	CCC Pro 55	GGC Gly	CGC Arg	CTT Leu	CCG Pro	AGC Ser 60	CTC Leu	CTT Leu	CCC Pro	CCC Pro			199
TCT Ser 65	CGA Arg	GAC Asp	GGG Gly	CCG Pro	GAA Glu 70	CCT Pro	CAA Gln	GAA Glu	GGA Gly	GGT Gly 75	CAG Gln	CCG Pro	GGT Gly	GCT Ala	GTG Val 80			247
CAA Gln	CGG Arg	TGC Cys	GCC Ala	TTG Leu 85	CCT Pro	CCC Pro	CGC Arg	TTG Leu	AAA Lys 90	GAG Glu	ATG Met	AAG Lys	AGT Ser	CAG Gln 95	GAG Glu			295
TCT Ser	GTG Val	GCA Ala	GGT Gly 100	TCC Ser	AAA Lys	CTA Leu	Val	CTT Leu 05	CGG Arg	TGC Cys	GAG Glu	Thr	AGT Ser 10	TCT Ser	GAA Glu			343
TAC Tyr	TCC Ser	TCT Ser 115	CTC Leu	AAG Lys	TTC Phe	Lys	TGG Trp 120	TTC Phe	AAG Lys	AAT Asn	GGG Gly	AGT Ser 125	GAA Glu	TTA Leu	AGC Ser			391
CGA Arg	AAG Lys 130	AAC Asn	AAA Lys	CCA Pro	Glu	AAC Asn 135	ATC . Ile	AAG Lys	ATA Ile	CAG Gln	AAA Lys 140	AGG Arg	CCG Pro	GGG Pro	AAG Lys			439
CA Ser 145	GAA Glu	CTT Leu	CGC Arg	ATT Ile	AGC Ser 150	AAA (Lys .	GCG Ala	TCA Ser	Leu	GCT Ala 155	GAT Asp	TCT Ser	GGA Gly	Glu	TAT Tyr 160			487

Figure 11 C'

ATG Met	TGC Cys	AAA Lys	GTG Val	ATC Ile 165	AGC Ser	AAA Lys	CTA Leu	GGA Gly	AAT Asn 170	GAC Asp	AGT Ser	GCC Ala	TCT Ser	GCC Ala 175	AAC Asn	535
ATC Ile	ACC Arg	ATT Ile	GTG Val 180	GAG Glu	TCA Ser	AAC Asn	GCC Ala	ACA Thr 185	TCC Ser	ACA Thr	TCT Ser	ACA Thr	GCT Ala 190	GGG Gly	ACA Thr	583
AGC Ser	CAT His	CTT Leu 195	GTC Val	AAG Lys	TGT Cys	Ala	GAG Glu 200	AAG Lys	GAG Glu	AAA Lys	ACT Thr	TTC Phe 205	TGT Cys	GTG Val	AAT Asn	631
GGA Gly	GGC Gly 210	GAG Glu	TGC Cys	TTC Phe	ATG Met	GTG Val 215	AAA Lys	GAC Asp	CTT Leu	TCA Ser	AAT <u>Asn</u> 220	CCC Pro	TCA Ser	AGA Arg	TAC Tyr	679
TTG Leu 225	TGC Cys	AAG Lys	TGC Cys	CCA Pro	AAT Asn 230	GAG Glu	TTT Phe	ACT Thr	GGT Gly	GAT Asp 235	CGC Arg	TGC Cys	CAA Gln	AAC Asn	TAC Tyr 240	727
GTA Val	ATG Met	GCC Ala	AGC Ser	TTC Phe 245	TAC Tyr	AGT Ser	ACG Thr	TCC Ser	ACT Thr 250	CCC Pro	TTT Phe	CTG Leu	TCT Ser	CTG Leu 255	CCT Pro	775
GAA Glu	TAG	CGCA'	rct (CAGT	CGGTY	GC C	GCTT	CTI	TT	GCCG	CATC	TCC	CCTC	AGA '	TTCCGCCTA	3 838
AGC'	raga:	rgc (GTTT:	racc.	AG G	ICTA	ACAT	r ga	CTGC	CTCT	GCC'	TGTC	GCA	TGAG.	AACATT	898
AAC	ACAA	GCG 2	ATTG:	ratg:	AC T	rccr	CTGT	C CG	rgac'	TAGT	GGG	CTCT	GAG	CTAC	TCGTAG	958
															TGATAG	1018
															AAAAA	1078
			GTCG													1108



17/52

Figure 13 A

CODING SEGMENT F:	
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GGCGGCTGCC CAGGCGATGC GAGCGCGGGC CGGACGGTAA TCGCCTCTCC CTCCTCGGGC	120
TGCGAGCGCG CCGGACCGAG GCAGCGACAG GAGCGGACCG CGGCGGGAAC CGAGGACTCC	180
CCAGCGGCGC GCCAGCAGGA GCCACCCCGC GAGNCGTGCG ACCGGGACGG AGCGCCCGCC	240
AGTCCCAGGT GGCCCGGACC GCACGTTGCG TCCCCGCGGCT CCCCGCCGGC GACAGGAGAC	300
GCTCCCCCC ACGCCGCGC CGCCTCGGCC CGGTCGCTGG CCCGCCTCCA CTCCGGGGAC	360
AAACTTTTCC CGAAGCCGAT CCCAGCCCTC GGACCCAAAC TTGTCGCGCG TCGCCTTCGC	420
CGGGAGCCGT CCGCGCAGAG CGTGCACTTC TCGGGCGAG ATG TCG GAG CGC AGA	474
Glu Gly Lys Gly Lys Gly Lys Gly Gly Lys Lys Asp Arg Gly Ser Gly GAA GGC AAA GGC AAG GGC GGC AAG AAG GAC CGA GGC TCC GGG	522
Lys Lys Pro Val Pro Ala Ala Gly Gly Pro Ser Pro Ala AAG AAG CCC GTG CCC GCG GCT GGC GGC CCG AGC CCA G	559

.,*

Figure 13 B

CODING	SEGMENT	E:	

CC	CAT His 1	CAN Gln	GTG Val	TGG Trp	GCG Ala 5	GCG Ala	AAA Lys	GCC Ala	GGG Gly	GGC Gly 10	TTG Leu	AAG Lys	AAG Lys	GAC Asp	TCG Ser 15	47
CTG Leu	CTC Leu	ACC Thr	GTG Val	CGC Arg 20	CTG Leu	GGC Gly	GCC Ala	TGG Trp	GGC Gly 25	CAC	Pro	GCC Ala	TTC Phe	CCC Pro	TCC Ser	95
TGC Cys	GGG	CGC Arg	CTC Leu 35	Lys	GAG Glu	GAC Asp	AGC Ser	AGG Arg 40	TAC Tyr	ATC Ile	TTC Phe	TTC Phe	ATG Met	Glu	CCC	143
GAG Glu	GCC Ala	AAC Asn 50	AGC Ser	AGC Ser	GGC Gly	GGG Gly	CCC Pro 55	GGC Gly	CGC Arg	CTT Leu	CCG Pro	AGC Ser 60	Leu	CTT Leu	CCC Pro	191
CCC Pro	TCT Ser 65	CGA Arg	GAC Asp	GGG Gly	CCG Pro	GAA Glu 70	CCT Pro	CAA Gln	GAA Glu	GGA Gly	GGT Gly 75	CAG Gln	CCG Pro	GGT Gly	GCT Ala	239
		CGG Arg		G							·					252

Figure 13 C

CODING SEGMENT B:

CCT	Pro TGC TGC	CTC	ccc	GCT	TGA	AAG	AGA	TGA	AGA	GTC	AGG	AGT 	CTG	TGG	CAG	48
GTT	Lys CCA CCA	AAC	TAG	TGC	TTC	GGT	GCG	AGA	CCA	GTT	CTG	AAT	ACT	CCT	CTC	96
TCA	Phe AGT GAT	TCA	AGT	GGT	TCA	AGA	ATG	GGA	GTG	AAT 	TAA	GCC	GAA	AGA	ACA	144
AAC	Gly CAC CAC	AAA 	ACA	TCA	AGA	TAC	AGA	AAA 	GGC	CGG	1					178

Figure 13 D

CODI	ING S	SEGMI	ent 1	A:												
GAA	G TO	CA GA	AA C	eu Ai FT CC	GC AT	PT AC	GC A/	AA GO	CG TY 	CA C'.	G G	.T G				46
GAA	TAT	ATG	TGC	Lys AAA AAA	GTG	ATC	AGC	AAA	CTA	GGA	AAT	III	AGI	111	iii	94
GCC	AAC	ATC	ACC	Ile ATT ATC	GTG 	GAG	TCA	AAC	G							122

Figure 13 E

CODING SEGMENT A':	
TCTAAAACTA CAGAGACTGT ATTTTCATGA TCATCATAGT TCTGTGAAAT ATACTTAAAC	60
CGCTTTGGTC CTGATCTTGT AGG AAG TCA GAA CTT CGC ATT AGC AAA GCG Lys Ser Glu Leu Arg Ile Ser Lys Ala 1	110
TCA CTG GCT GAT TCT GGA GAA TAT ATG TGC AAA GTG ATC AGC AAA CTA Ser Leu Ala Asp Ser Gly Glu Tyr Met Cys Lys Val Ile Ser Lys Leu 10 25	158
GGA AAT GAC AGT GCC TCT GCC AAC ATC ACC ATT GTG GAG TCA AAC GGT Gly Asn Asp Ser Ala Ser Ala Asn Ile Thr Ile Val Glu Ser Asn Gly 30 35 40	206
AAG AGA TGC CTA CTG CGT GCT ATT TCT CAG TCT CTA AGA GGA GTG ATC Lys Arg Cys Leu Leu Arg Ala Ile Ser Gln Ser Leu Arg Gly Val Ile 45 50 55	254
AAG GTA TGT GGT CAC ACT TGAATCACGC AGGTGTGTGA AATCTCATTG Lys Val Cys Gly His Thr 60	302
TGAACAAATA AAAATCATGA AAGGAAAACT CTATGTTTGA AATATCTTAT GGGTCCTCCT	362
GTAAAGCTCT TCACTCCATA AGGTGAAATA GACCTGAAAT ATATATAGAT TATTT	417

Figure 13 F

CODING	SEGMENT	G:
--------	---------	----

II	III	ΙĬ	Thr ACT ACT I	11	III	III	111	ICA	ACT	GAG	ACA	GCG	TAT	GTG		47
111	III	iii	Pro CCC CCC		AGA	III	ICA	GTA	TCA	ACA	GAA	GGA	ACA	AAT	ACT	95
Ser TCT TCT	TCA	T											A			102

Figure 13 G

CODING SEGMENT C:

Thr Ser Thr Ser Thr Ala Gly Thr Ser His Leu Val Lys Cys Ala CC ACA TCC ACA TCT ACA GCT GGG ACA AGC CAT CTT GTC AAG TGT GCA	47
Glu Lys Glu Lys Thr Phe Cys Val Asn Gly Gly Glu Cys Phe Met Val GAG AAG GAG AAA ACT TTC TGT GTG AAT GGA GGC GAG TGC TTC ATG GTG	95
Lys Asp Leu Ser Asn Pro Ser Arg Tyr Leu Cys AAA GAC CTT TCA AAT CCC TCA AGA TAC TTG TGC AAA GAC CTT TCA AAC CCC TCG AGA TAC TTG TGC	128

Figure 13 H

CODING SEGMENT C/D:	
---------------------	--

Lys AAG AAG	TGC	CAA	CCT	GGA	TTC	ACT	GGA	II	III	111	III	III	ĬĬĬ	ĬĬĬ	41	В
ATG	AAA	GTC	Gln CAA CAA	ACC	CAA	GAA									6	9

Figure 13 I

CODING SEGMENT C/D':

	TGC	CCA	AAT	GAG	TTT	ACT	GGT	III	111	111	111	III	111	Met ATG ATG	48
Ala GCC GCC	AGC	TTC	TAC												60

Figure 13 J

CODING SEGMENT D:

			Thr ACT ACT	~~~	THE STATE OF		יויייאווי	CHY	C.TI.	LiAA	TAG		3
--	--	--	--------------------------	-----	--------------	--	-----------	-----	-------	------	-----	--	---

Figure 13 K

CODING SEGMENT D':

Lys His Leu Gly Ile Glu Phe Met Glu AAG CAT CTT GGG ATT GAA TTT ATG GAG	27
--	----

Figure 13 L

CODING SEGMENT H:

Val Leu Thr Ile Thr Gly Ile GTG CTC ACC ATT ACC GGC ATT GTG CTG ACC ATA ACC GGC ATC	48
Met Cys Val Val Val Tyr Cys ATG TGT GTG GTG GTC TAC TGC ATG TGT GTG GTG GCC TAC TGC A	96
His Asp Arg Leu Arg Gln Ser CAT GAC CGG CTT CGG CAG AGC CAT GAC CGT CTT CGG CAG AGC	144
Asn Val Ala Asn Gly Pro His AAC GTA GCC AAC GGG CCC CAC AAC ATT GCC AAT GGG CCT CAC	192
Gln Leu Val Asn Gln Tyr Val CAG CTG GTG AAT CAA TAC GTA CAG CTG GTG AAT CAA TAC GTA	2 4 0
Ile Val Glu Arg Glu Ala Glu ATT GTT GAG AGA GAG GCG GAG ATT GTT GAG AGA GAA GCA GAG	288

Figure 13 L'

AGC	TCT	Phe TTT TTT	TCC	ACC	AGT	CAC	TAC	ACT	TCG	ACA	GCT 		LAT	111	III	336
ACT	GTC	Thr ACT ACC	CAG	ACT	CCC	AGT	CAC	AGC	TGG	AGC	AAT	GGA	III	ACT	GAA	384
AGC	ATC	Ile ATT CTT L	TCG	GAA	AGC	CAC	TCT	GTC	ATC		AIG	III	111			432
AAC	AGT	Arg AGG AGG	CAC	AGC	AGC	CCG	ACT	GGG	GGC	CCG	AGA	GGA		11	HAT	480
GGC	TTG	Gly GGA GGA	GGC	CCT	CGT	GAA	TGT	AAC	AGC	III	111	AGG	III	111	III	528
GAA	ACC	Pro CCT	GAC	TCC	TAC	CGA	GAC	TCT	CCT	CAT	AGT	III	AG 			569

WO 96/15812 PCT/US95/14974

Figure 13 M

CODING SEGMENT K:

A	CAT His . 1	AAC Asn	Leu	II	A G e A	CT G la G 5	AG C lu L	TA A eu A	GG A rg A	GA A rg A	AC A sn L 10	AG G ys A	CC C la H	AC A	GA T	CC er 15	46
AA! Lys	TGG Cy:	C AT s Me	t G	AG .	ATC Ile 20	CAG Gln	CTT Leu	TCC Ser	GCA Ala	ACT Thr 25	CAT His	CTT Leu	AGA Arg	GCT Ala	TCT Ser 30	TCC Ser	94
ATT Ile	CCC Pro	C CA Hi	T T s T	GG (rp 2 35	GCT Ala	TCA Ser	TTC Phe	TCT Ser	AAG Lys 40	ACC Thr	CCT Pro	TGG Trp	CCT Pro	TTA Leu 45	GGA Gly	AG Arg	141

Figure 13 N

CODING SEGMENT L:

Tyr Val Ser Ala Met Thr Thr Pro Ala Arg Met Ser Pro Val Asp G TAT GTA TCA GCA ATG ACC ACC CCG GCT CGT ATG TCA CCT GTA GAT	46
Phe His Thr Pro Ser Ser Pro Lys Ser Pro Pro Ser Glu Met Ser Pro TTC CAC ACG CCA AGC TCC CCC AAG TCA CCC CCT TCG GAA ATG TCC CCG	94
Pro Val Ser Ser Thr Thr Val Ser Met Pro Ser Met Ala Val Ser Pro CCC GTG TCC AGC ACG GTC TCC ATG CCC TCC ATG GCG GTC AGT CCC	142
Phe Val Glu Glu Glu Arg Pro Leu Leu Val Thr Pro Pro Arg Leu TTC GTG GAA GAG GAG AGA CCC CTG CTC CTT GTG ACG CCA CCA CGG CTG	190
Arg Glu Lys - Tyr Asp His His Ala Gln Gln Phe Asn Ser Phe His CGG GAG AAG TAT GAC CAC CAC GCC CAG CAA TTC AAC TCG TTC CAC	238
Cys Asn Pro Ala His Glu Ser Asn Ser Leu Pro Pro Ser Pro Leu Arg TGC AAC CCC GCG CAT GAG AGC AAC AGC CTG CCC CCC AGC CCC TTG AGG	286

Figure 13 N'

ATA	Val GTG GTG	GAG	GAT 	GAG	GAA 	TAT	GAA	ACG	ACC	CAG	GAG	TAC	GAA	CCA	Ala GCT GCC	334
CAA	Glu GAG GAG	CCG	GTT	AAG	AAA 	CTC	ACC	AAC	AGC	AGC	CGG	CGG	GCC 	AAA	AGA	382
ACC	Lys AAG AAG	ccc	TAA	GGT	CAC	ATT	GCC	CAC	AGG	TTG	GAA	ATG	GAC	AAC 	AAC 	430
Thr ACA ACA	-	GCT TCC	GAC	AGC	AGT	AAC	TCA	GAG	AGC	GAA	ACA	GAG	GAT	GAA	AGA	478

Figure 13 N''

GTA	GGA	Glu GAA GAA	GAT	ACG	CCT	TTC	CIG	GCC	III	III		III	ĬĬĬ	ĬII	ĬĬĬ	52	6
AGT	CTC	Glu GAG GAG	ĢÇG	GCC	CCT	GUU	111	111	111	J.C	111	111	111	TIL	111	57	4
CCA CCA	ACA		GĢÇ	TTC	TCT	CCG	LAG	III	III	1	111	III	111	ĪĪ	Ser TCC TCT	62	<u>?</u> 2
GGT AGT S	GTA GTA	ATT	GCT GCT	AAC AAC	CAA	GAC GAC	CCT	ATT	GCT	 GTA	TAA	AAC	CTA	AAT	ACA AAA		72
															CCA CCA		18
111	111	ATT ATT		-111												7:	33

Figure 13 0

HUMAN CODING SEGMENT E:

ATG Met 1	AG Arg	A TG g Tr	G CG p Ar	A CG g Ar 5	C GC g Al	C CC a Pr	G CG o Ar	C CG g Ar	TC Se:	C GGG r Gl	G CG Y Ar	r CCC	GGC Gly	C CCC 7 Pro 15	C CGG Arg	• 48
GCC Ala	CAC Glr	G CG	C CC g Pr 20	o Gl	C TC y Se	C GC r Al	C GCG a Ala	C CGG Arg 25	TC(g Sei	TC(G CCC	G CCC	CTC Leu 30	CCC Pro	CTG Leu	96
CTG Leu	Pro	CT. Let	A CTO	G CT u Le	G CT	G CT u Le	G GG(1 Gl) 40	G ACC	GCC Ala	G GCC	CTY Leu	GCG Ala 45	CCG Pro	GGG Gly	GCG Ala	144
GCG Ala	GCC Ala 50	GGG Gly	AAC Ası	C GAG	G GCG	G GC' a Ala 55	r ccc	GCG Ala	GGG Gly	GCC Ala	TCC Ser 60	GTG Val	TGC Cys	TAC Tyr	TCG Ser	192
TCC Ser 65	CCG Pro	Pro	AGC Ser	C GT(GG/ Gl ₃ 70	A TCC	GTG Val	CAG Gln	GAG Glu	CTA Leu 75	GCT Ala	CAG Gln	CGC Arg	GCC Ala	GCG Ala 80	240
GTG Val	GTG Val	ATC	GAG Glu	GGA Gly 85	AAC Lys	GTC Val	CAC His	CCG Pro	CAG Gln 90	CGG Arg	CGG Arg	CAG Gln	CAG Gln	GGG Gly 95	GCA Ala	288
CTC Leu	GAC Asp	AGG Arg	AAG Lys 100	Ala	GCG	GCG Ala	Ala	GCG Ala 105	GGC Gly	GAG Glu	GCA Ala	GGG Gly	GCG Ala 110	TGG Trp	GGC GLY	336
GGC	GAT Asp	CGC Arg 115	Glu	CCG Pro	CCA Pro	GCC Ala	GCG Ala 120	GGC Gly	CCA Pro	CGG Arg	GCG Ala	CTG Leu 125	GGG Gly	CCG Pro	CCC Pro	384
GCC Ala	GAG Glu 130	GAG Glu	CCG Pro	CTG Leu	CTC Leu	GCC Ala 135	GCC Ala	AAC Asn	GGG Gly	ACC Thr	GTG Val 140	CCC Pro	TCT Ser	TGG Trp	CCC Pro	432
ACC (Thr 145	GCC Ala	CCG Pro	GTG Val	CCC Pro	AGC Ser 150	GCC Ala	GGC Gly	GAG Glu	CCC Pro	GGG Gly 155	GAG Glu	GAG Glu	GCG Ala	CCC Pro	TAT Tyr 160	480
CTG (Leu 1	GTG Val	AAG Lys	GTG Val	CAC His 165	CAG Gln	GTG Val	TGG Trp	GCG Ala	GTG Val 170	AAA Lys	GCC Ala	GGG Gly	GGC Gly	TTG Leu 175	AAG Lys	528
AAG (Lys 1	GAC Asp	TCG Ser	CTG Leu 180	CTC Leu	ACC Thr	GTG Val	CGC Arg	CTG Leu 185	GGG Gly	ACC Thr	TGG Trp	Gly	CAC His 190	CCC Pro	GCC Ala	576
TTC C	Pro	TCC Ser 195	TGC Cys	GGG Gly	AGG Arg	CTC Leu	AAG Lys 200	GAG Glu	GAC Asp	AGC Ser	AGG Arg	TAC Tyr 205	ATC Ile	TTC Phe	TTC Phe	624
ATG G Met G 2	AG (lu l	CCC Pro	GAC Asp	GCC Ala	AAC Asn	AGC Ser 215	ACC Thr	AGC Ser	CGC Arg	Ala	CCG Pro 220	GCC Ala	GCC Ala	TTC Phe	CGA Arg	672
GCC T Ala S 225	CT :	MC Phe	CCC Pro	Pro :	CTG Leu 230	GAG Glu	ACG (Thr	GGC (Gly)	Arg A	AAC Asn 235	CTC Leu	AAG . Lys :	AAG (Lys (Glu	GTC Val 240	720
AGC C Ser A	GG 0 rg V	TG (Leu (TGC Z Cys 1 245	AAG (Lys)	CGG Arg (IGC (Cys	3	207	5 0						745

Figure 14 A

AGTTTCCCCC CCCAACTTGT CGGAACTCTG GGCTCGCGCG CAGGGCAGGA GCGGAGCGGC	60
GGCGGCTGCC CAGGCGATGC GAGCGCGGGC CGGACGGTAA TCGCCTCTCC CTCCTCGGGC	120
TGCGAGCGCG CCGGACCGAG GCAGCGACAG GAGCGGACCG CGGCGGGAAC CGAGGACTCC	180
CCAGCGGCGC GCCAGCAGGA GCCACCCCGC GAGCGTGCGA CCGGGACGGA GCGCCCGCCA	240
GTCCCAGGTG GCCCGGACCG CACGTTGCGT CCCCGCGCTC CCCGCCGGCG ACAGGAGACG	300
CTCCCCCCA CGCCGCGC GCCTCGGCCC GGTCGCTGGC CCGCCTCCAC TCCGGGGACA	360
AACTTTTCCC GAAGCCGATC CCAGCCCTCG GACCCAAACT TGTCGCGCGT CGCCTTCGCC	420
GGGAGCCGTC CGCGCAGAGC GTGCACTTCT CGGGCGAG ATG TCG GAG CGC AGA Met Ser Glu Arg Arg 1 5	475
GAA GGC AAA GGC AAG GGG AAG GGC GGC AAG AAG	523
AAG AAG CCC GTG CCC GCG GCT GGC GGC CCG AGC CCA GCC TTG CCT CCC Lys Lys Pro Val Pro Ala Ala Gly Gly Pro Ser Pro Ala Leu Pro Pro 25 30 35	571
CGC TTG AAA GAG ATG AAG ATG CAG GAG TCT GTG GCA GGT TCC AAA CTA Arg Leu Lys Glu Met Lys Ser Gln Glu Ser Val Ala Gly Ser Lys Leu 40 45	619
GTG CTT CGG TGC GAG ACC AGT TCT GAA TAC TCC TCT CTC AAG TTC AAG Val Leu Arg Cys Glu Thr Ser Ser Glu Tyr Ser Ser Leu Lys Phe Lys 55 60 65	667
TGG TTC AAG AAT GGG AGT GAA TTA AGC CGA AAG AAC AAA CCA CAA AAC Trp Phe Lys Asn Gly Ser Glu Leu Ser Arg Lys Asn Lys Pro Gln Asn 70 80 85	715
ATC AAG ATA CAG AAA AGG CCG GGG AAG TCA GAA CTT CGC ATT AGC AAA Ile Lys Ile Gln Lys Arg Pro Gly Lys Ser Glu Leu Arg Ile Ser Lys 90 95 100	763
GCG TCA CTG GCT GAT TCT GGA GAA TAT ATG TGC AAA GTG ATC AGC AAA ALA Ser Leu Ala Asp Ser Gly Glu Tyr Met Cys Lys Val Ile Ser Lys 105	811

Figure 14 B

CTA GGA AAT GAC AGT GCC TCT GCC AAC ATC ACC ATT GTG GAG TCA AAC Leu Gly Asn Asp Ser Ala Ser Ala Asn Ile Thr Ile Val Glu Ser Asn 120 125 130	859
GAG ATC ACC ACT GGC ATG CCA GCC TCA ACT GAG ACA GCG TAT GTG TCT Glu Ile Thr Thr Gly Met Pro Ala Ser Thr Glu Thr Ala Tyr Val Ser 135 140 145	907
TCA GAG TCT CCC ATT AGA ATA TCA GTA TCA ACA GAA GGA ACA AAT ACT Ser Glu Ser Pro Ile Arg Ile Ser Val Ser Thr Glu Gly Thr Asn Thr 150 165	955
TCT TCA TCC ACA TCC ACA TCT ACA GCT GGG ACA AGC CAT CTT GTC AAG Ser Ser Ser Thr Ser Thr Ala Gly Thr Ser His Leu Val Lys 170 175 180	1003
TGT GCA GAG AAG GAG AAA ACT TTC TGT GTG AAT GGA GGC GAG TGC TTC Cys Ala Glu Lys Glu Lys Thr Phe Cys Val Asn Gly Gly Glu Cys Phe 185 190 195	1051
ATG GTG AAA GAC CTT TCA AAT CCC TCA AGA TAC TTG TGC AAG TGC CCA Met Val Lys Asp Leu Ser Asn Pro Ser Arg Tyr Leu Cys Lys Cys Pro 200 205 210	1099
AAT GAG TTT ACT GGT GAT CGC TGC CAA AAC TAC GTA ATG GCC AGC TTC Asn Glu Phe Thr Gly Asp Arg Cys Gln Asn Tyr Val Met Ala Ser Phe 215 220 225	1147
TAC AGT ACG TCC ACT CCC TTT CTG TCT CTG CCT GAA TAGGCGCATG Tyr Ser Thr Ser Thr Pro Phe Leu Ser Leu Pro Glu 230 240	1193
CTCAGTCGGT GCCGCTTTCT TGTTGCCGCA TCTCCCCTCA GATTCAACCT AGAGCTAGAT	1253
GCGTTTTACC AGGTCTAACA TTGACTGCCT CTGCCTGTCG CATGAGAACA TTAACACAAG	1313
CGATTGTATG ACTTCCTCTG TCCGTGACTA GTGGGCTCTG AGCTACTCGT AGGTGCGTAA	1373
GGCTCCAGTG TTTCTGAAAT TGATCTTGAA TTACTGTGAT ACGACATGAT AGTCCCTCTC	1433
ACCCAGTGCA ATGACAATAA AGGCCTTGAA AAGTCTCACT TTTATTGAGA AAATAAAAAT	1493
CGTTCCACGG GACAGTCCCT CTTCTTTATA AAATGACCCT ATCCTTGAAA AGGAGGTGTG	1553
TTAAGTTGTA ACCAGTACAC ACTTGAAATG ATGGTAAGTT CGCTTCGGTT CAGAATGTGT	1613
TCTTTCTGAC AAATAAACAG AATAAAAAAA AAAAAAAAAA	1654

Figure 15 A

CAT His 1	CAN Gln	GTG Val	TGG Trp	GCG Ala 5	GCG Ala	AAA Lys	GCC Ala	GGG Gly	GGC Gly 10	TTG Leu	AAG Lys	AAG Lys	GAC Asp	TCG Ser 15	CTG Leu	48
CTC Leu	ACC Thr	GTG Val	CGC Arg 20	CTG Leu	GGC Gly	GCC Ala	TGG Trp	GGC Gly 25	CAC His	CCC Pro	GCC Ala	TTC Phe	CCC Pro 30	TCC Ser	TGC Cys	96
GGG Gly	CGC Arg	CTC Leu 35	AAG Lys	GAG Glu	GAC Asp	AGC Ser	AGG Arg 40	TAC Tyr	ATC Ile	TTC Phe	TTC Phe	ATG Met 45	GAG Glu	CCC Pro	GAG Glu	144
GCC Ala	AAC Asn 50	AGC Ser	AGC Ser	GGC Gly	GGG Gly	CCC Pro 55	GGC Gly	CGC Arg	CTT Leu	CCG Pro	AGC Ser 60	CTC Leu	CTT Leu	CCC Pro	CCC Pro	192
TCT Ser 65	CGA Arg	GAC Asp	GGG Gly	CCG Pro	GAA Glu 70	CCT Pro	CAA Gln	GAA Glu	GGA Gly	GGT Gly 75	CAG Gln	CCG Pro	GGT Gly	GCT Ala	GTG Val 80	240
CAA Gln	CGG Arg	TGC Cys	GCC Ala	TTG Leu 85	CCT Pro	CCC Pro	CGC Arg	TTG Leu	AAA Lys 90	GAG Glu	ATG Met	AAG Lys	AGT Ser	CAG Gln 95	GAG Glu	288
TCT Ser	GTG Val	GCA Ala	GGT Gly 100	TCC Ser	AAA Lys	CTA Leu	GTG Val	CTT Leu 105	CGG Arg	TGC Cys	GAG Glu	ACC Thr	AGT Ser 110	TCT Ser	GAA Glu	336
TAC Tyr	TCC Ser	TCT Ser 115	CTC Leu	AAG Lys	TTC Phe	AAG Lys	TGG Trp 120	TTC Phe	AAG Lys	AAT Asn	GGG Gly	AGT Ser 125	GAA Glu	TTA Leu	AGC Ser	384
CGA Arg	AAG Lys 130	AAC Asn	AAA Lys	CCA Pro	GAA Glu	AAC Asn 135	ATC Ile	AAG Lys	ATA Ile	CAG Gln	AAA Lys 140	AGG Arg	CCG Pro	GGG Gly	AAG Lys	432
TCA Ser 145	GAA Glu	CTT Leu	CGC Arg	ATT Ile	AGC Ser 150	AAA Lys	GCG Ala	TCA Ser	CTG Leu	GCT Ala 155	GAT Asp	TCT Ser	GGA Gly	GAA Glu	TAT Tyr 160	480
ATG Met	TGC Cys	AAA Lys	GTG Val	ATC Ile 165	AGC Ser	AAA Lys	CTA Leu	GGA Gly	AAT Asn 170	Asp	AGT Ser	GCC Ala	TCT Ser	GCC Ala 175	AAC Asn	528

Figure 15 B

Ile Thr Ile Val Glu Ser Asn Ala Thr Ser Thr Ser Thr Ala Gly Thr 180 185	576
AGC CAT CTT GTC AAG TGT GCA GAG AAG GAG AAA ACT TTC TGT GTG AAT Ser His Leu Val Lys Cys Ala Glu Lys Glu Lys Thr Phe Cys Val Asn 195 200 205	. 624
GGA GGC GAG TGC TTC ATG GTG AAA GAC CTT TCA AAT CCC TCA AGA TAC Gly Gly Glu Cys Phe Met Val Lys Asp Leu Ser Asn Pro Ser Arg Tyr 210 220	672
TTG TGC AAG TGC CAA CCT GGA TTC ACT GGA GCG AGA TGT ACT GAG AAT Leu Cys Lys Cys Gln Pro Gly Phe Thr Gly Ala Arg Cys Thr Glu Asn 230 235 240	720
GTG CCC ATG AAA GTC CAA ACC CAA GAA AAG TGC CCA AAT GAG TTT ACT Val Pro Met Lys Val Gln Thr Gln Glu Lys Cys Pro Asn Glu Phe Thr 245 250 255	768
GGT GAT CGC TGC CAA AAC TAC GTA ATG GCC AGC TTC TAC AGT ACG TCC Gly Asp Arg Cys Gln Asn Tyr Val Met Ala Ser Phe Tyr Ser Thr Ser 260 270	816
ACT CCC TTT CTG TCT CTG CCT GAA TAGCGCATCT CAGTCGGTGC CGCTTTCTTG Thr Pro Phe Leu Ser Leu Pro Glu 275 280	870
TTGCCGCATC TCCCCTCAGA TTCCNCCTAG AGCTAGATGC GTTTTACCAG GTCTAACATT	930
GACTGCCTCT GCCTGTCGCA TGAGAACATT AACACAAGCG ATTGTATGAC TTCCTCTGTC	990
CGTGACTAGT GGGCTCTGAG CTACTCGTAG GTGCGTAAGG CTCCAGTGTT TCTGAAATTG	1050
ATCTTGAATT ACTGTGATAC GACATGATAG TCCCTCTCAC CCAGTGCAAT GACAATAAAG	1110
GCCTTGAAAA GTCAAAAAAA AAAAAAAAA	1140

Figure 16 A

G AA Ly 1	's Se	A GA r Gl	A CT u Le	T CG u Ar 5	C AT	T AG e Se	C AA r Ly	A GC	a cc	A CT r Le 0	G GC u Al	T GA' a As	T TC	T GG r Gl	A GAA y Glu 5	49
		TGC Cys	AAA Lys 20	GTG Val	ATC Ile	AGC Ser	AAA Lys	CTA Leu 25	GGA Gly	AAT Asn	GAC Asp	AGT Ser	GCC Ala 30	TCT Ser	GCC Ala	97
AAC Asn	ATC Ile	ACC Thr 35	ATT Ile	GTG Val	GAG Glu	TCA Ser	AAC Asn 40	GCC Ala	ACA Thr	TCC Ser	ACA Thr	TCT Ser 45	ACA Thr	GCT Ala	GGG Gly	145
ACA Thr	AGC Ser 50	CAT His	CTT Leu	GTC Val	AAG Lys	TGT Cys 55	GCA Ala	GAG Glu	AAG Lys	GAG Glu	AAA Lys 60	ACT Thr	TTC Phe	TGT Cys	GTG Val	193
AAT Asn 65	GGA Gly	GGC Gly	GAC Asp	TGC Cys	TTC Phe 70	ATG Met	GTG Val	AAA Lys	GAC Asp	CTT Leu 75	TCA Ser	AAT Asn	CCC Pro	TCA Ser	AGA Arg 80	241
TAC Tyr	TTG Leu	TGC Cys	AAG Lys	TGC Cys 85	CAA Gln	CCT Pro	GGA Gly	TTC Phe	ACT Thr 90	GGA Gly	GCG Ala	AGA Arg	TGT Cys	ACT Thr 95	GAG Glu	289
AAT Asn	GTG Val	CCC Pro	ATG Met 100	AAA Lys	GTC Val	CAA Gln	TIII	CAA Gln 105	GAA Glu	AAA Lys	GCG Ala	GAG Glu	GAG Glu 110	CTC Leu	TAC Tyr	337
CAG Gln	AAG Lys	AGA Arg 115	Val	CTC Leu	ACC Thr	ATT Ile	ACC Thr 120	GGC Gly	ATT Ile	TGC Cys	ATC Ile	GCG Ala 125	CTG Leu	CTC Leu	GTG Val	385
GTT Val	GGC Gly 130	Ile	ATG Met	TGT Cys	GTG Val	GTG Val 135	GTC Val	TAC Tyr	TGC Cys	AAA Lys	ACC Thr 140	AAG Lys	AAA Lys	CAA Gln	CGG Arg	433
Lys 145	Lys	Leu	His	Asp	150	Leu	Arg	GII	Ser	155	2				AAC Asn 160	481
Thr	Met	Met	Asn	165	Ala	ASI	GIZ	PIC	170)				175	CCC Pro	529
GAG Glu	AAC Asr	GTG Val	CAG Gln 180	Leu	GTG Val	AAT Asn	CAA Gln	TAC Ty: 185	. VO.	TCI Ser	Lys	AAT AST	Val 190	ATC Ile	Ser	577

Figure 16 B

AG Se:	GA(G CA Hi 19	s II	T GT e Va	r gad	G AGA u Arg	GAG Glu 200	ı Ala	G GAG	G AGG	C TC	r TT r Pho 20	e Se	C AC	C AGT r Ser		625
CA(His	TAC 5 Tyr 210	rTh	T TC	G ACI	A GCT	CAT His 215	His	TCC Ser	Thi	Thi	r GT(Va. 220	l Thi	r Gli	G AC	r CCC		673
AGT Ser 225	Hls	S Sei	C TG(G AGO Ser	AAT Asr 230	ı Gly	CAC His	ACT Thr	GAA	AGO Ser 235	: Ile	ATT	TCC Sei	GAA	A AGC Ser 240		721
CAC His	TCI Ser	GTC Val	Ile	GTG Val 245	Met	TCA Ser	TCC Ser	GTA Val	GAA Glu 250	Asn	AG1 Ser	AGG Arg	CAC His	AGC Ser 259	AGC Ser		769
CCG Pro	ACT	GGG	GGC Gly 260	Pro	AGA Arg	GGA Gly	CGT Arg	CTC Leu 265	AAT Asn	GGC Gly	TTG	GGA Gly	GGC Gly 270	Pro	CGT Arg		817
GAA Glu	TGT Cys	AAC Asn 275	Ser	TTC Phe	CTC Leu	AGG Arg	CAT His 280	GCC Ala	AGA Arg	GAA Glu	ACC Thr	CCT Pro 285	GAC Asp	TCC	TAC Tyr		865
CGA Arg	GAC Asp 290	TCT Ser	CCT Pro	CAT His	AGT Ser	GAA Glu 295	AGA Arg	CAT His	AAC Asn	CTT Leu	ATA Ile 300	GCT Ala	GAG Glu	CTA Leu	AGG Arg		913
AGA Arg 305	AAC Asn	AAG Lys	GCC Ala	CAC His	AGA Arg 310	TCC Ser	AAA Lys	TGC Cys	ATG Met	CAG Gln 315	ATC Ile	CAG Gln	CTT Leu	TCC Ser	GCA Ala 320		961
ACT Thr	CAT His	CTT Leu	AGA Arg	GCT Ala 325	TCT Ser	TCC Ser	ATT Ile	CCC Pro	CAT His 330	TGG Trp	GCT Ala	TCA Ser	TTC Phe	TCT Ser 335	AAG Lys		1009
ACC Thr	CCT Pro	TGG Trp	CCT Pro 340	TTA Leu	GGA Gly	AGG Arg	Tyr	GTA Val 345	TCA Ser	GCA Ala	ATG Met	ACC Thr	ACC Thr 350	CCG Pro	GCT Ala		1057
CGT Arg	Met	TCA Ser 355	CCT Pro	GTA Val	GAT Asp	TTČ Phe	CAC A His '	ACG Thr	CCA Pro	AGC Ser	TCC Ser	CCC Pro 365	AAG Lys	TCA Ser	CCC Pro	:	1105
Pro	TCG Ser 370	GAA Glu	ATG Met	TCC Ser	Pro	CCC (Pro 1 375	GTG : Val :	CC . Ser	AGC Ser	Thr	ACG Thr 380	GTC Val	TCC Ser	ATG Met	CCC Pro	;	1153

Figure 16 C

TCC ATG GCG GTC AGT CCC TTC GTG GAA GAG GAG AGA CCC CTG C Ser Met Ala Val Ser Pro Phe Val Glu Glu Arg Pro Leu L 385 390 395	CTC CTT 1201 Leu Leu 400
GTG ACG CCA CGG CTG CGG GAG AAG TAT GAC CAC CAC GCC CVal Thr Pro Pro Arg Leu Arg Glu Lys Tyr Asp His His Ala G	CAG CAA 1249 Gln Gln 115
TTC AAC TCG TTC CAC TGC AAC CCC GCG CAT GAG AGC AAC AGC C Phe Asn Ser Phe His Cys Asn Pro Ala His Glu Ser Asn Ser L 420 425 430	CTG CCC 1297 Leu Pro
CCC AGC CCC TTG AGG ATA GTG GAG GAT GAG GAA TAT GAA ACG A Pro Ser Pro Leu Arg Ile Val Glu Asp Glu Glu Tyr Glu Thr T 435	ACC CAG 1345 Thr Gln
GAG TAC GAA CCA GCT CAA GAG CCG GTT AAG AAA CTC ACC AAC AG Glu Tyr Glu Pro Ala Glu Pro Val Lys Lys Leu Thr Asn S 450	AGC AGC 1393 Ser Ser
CGG CGG GCC AAA AGA ACC AAG CCC AAT GGT CAC ATT GCC CAC AATG ATG ATG Ala Lys ATG Thr Lys Pro Asn Gly His Ile Ala His AA65 470 475	AGG TTG 1441 Arg Leu 480
GAA ATG GAC AAC AAC ACA GGC GCT GAC AGC AGT AAC TCA GAG AGU Met Asp Asn Asn Thr Gly Ala Asp Ser Ser Asn Ser Glu S	AGC GAA 1489 Ser Glu 495
ACA GAG GAT GAA AGA GTA GGA GAA GAT ACG CCT TTC CTG GCC A Thr Glu Asp Glu Arg Val Gly Glu Asp Thr Pro Phe Leu Ala I 500 505 510	ATA CAG 1537 Ile Gln
AAC CCC CTG GCA GCC AGT CTC GAG GCG GCC CCT GCC TTC CGC CAsn Pro Leu Ala Ala Ser Leu Glu Ala Ala Pro Ala Phe Arg I 515	CTG GTC 1585 Leu Val
GAC AGC AGG ACT AAC CCA ACA GGC GGC TTC TCT CCG CAG GAA CA ASP Ser Arg Thr Asn Pro Thr Gly Gly Phe Ser Pro Gln Glu G 530 540	GAA TTG 1633 Glu Leu
CAG GCC AGG CTC TCC GGT GTA ATC GCT AAC CAA GAC CCT ATC Gln Ala Arg Leu Ser Gly Val Ile Ala Asn Gln Asp Pro Ile A	GCT GTC 1681 Ala Val 560
TAAAACCGAA ATACACCCAT AGATTCACCT GTAAAACTIT ATITTATATA AT	TAAAGTATT 1741
CCACCTTAAA TTAAACAAAA AAA	1764

Figure 17 A

F-B-A'

F-E-B-A'

F-B-A-C-C/D-D F-E-B-A-C-C/D-D F-B-A-C-C/D-H F-E-B-A-C-C/D-H F-B-A-C-C/D-H-L F-E-B-A-C-C/D-H-L F-B-A-C-C/D-H-K-L F-E-B-A-C-C/D-H-K-L F-B-A-C-C/D-D'-H F-E-B-A-C-C/D-D'-H F-B-A-C-C/D-D'-H-L F-E-B-A-C-C/D-D'-H-L F-B-A-C-C/D-D'-H-K-L F-E-B-A-C-C/D-D'-H-K-L F-B-A-C-C/D'-D F-E-B-A-C-C/D'-D F-B-A-C-C/D'-H F-E-B-A-C-C/D'-H F-B-A-C-C/D'-H-L F-E-B-A-C-C/D'-H-L F-B-A-C-C/D'-H-K-L F-E-B-A-C-C/D'-H-K-L F-B-A-C-C/D'-D'-H F-E-B-A-C-C/D'-D'-H F-B-A-C-C/D'-D'-H-L F-E-B-A-C-C/D'-D'-H-L F-B-A-C-C/D'-D'-'H-K-L F-E-B-A-C-C/D'-D'-'H-K-L F-B-A-C-C/D-C/D'-D F-E-B-A-C-C/D-C/D'-D F-B-A-C-C/D-C/D'-H F-E-B-A-C-C/D-C/D'-H F-B-A-C-C/D-C/D'-H-L F-E-B-A-C-C/D-C/D'-H-L F-B-A-C-C/D-C/D'-H-K-L F-E-B-A-C-C/D-C/D'-H-K-L F-B-A-C-C/D-C/D'-D'-H F-E-B-A-C-C/D-C/D'-D'-H F-B-A-C-C/D-C/D'-D'-H-L F-E-B-A-C-C/D-C/D'-D'-H-L F-B-A-C-C/D-C/D'-D'-H-K-L F-E-B-A-C-C/D-C/D'-D'-H-K-L

F-B-A-G-C-C/D-D F-B-A-G-C-C/D-H F-B-A-G-C-C/D-H-L F-B-A-G-C-C/D-H-K-L F-B-A-G-C-C/D-D'-H F-B-A-G-C-C/D-D'-H-L F-B-A-G-C-C/D-D'-H-K-L F-B-A-G-C-C/D'-D F-B-A-G-C-C/D'-H F-B-A-G-C-C/D'-H-L F-B-A-G-C-C/D'-H-K-L F-B-A-G-C-C/D'-D'-H F-B-A-G-C-C/D'-D'-H-L F-B-A-G-C-C/D'-D'-'H-K-L F-B-A-G-C-C/D-C/D'-D F-B-A-G-C-C/D-C/D'-H F-B-A-G-C-C/D-C/D'-H-L F-B-A-G-C-C/D-C/D'-H-K-L F-B-A-G-C-C/D-C/D'-D'-H F-B-A-G-C-C/D-C/D'-D'-H-L F-B-A-G-C-C/D-C/D'-D'-H-K-L

F-E-B-A-G-C-C/D-D F-E-B-A-G-C-C/D-H F-E-B-A-G-C-C/D-H-L F-E-B-A-G-C-C/D-H-K-L F-E-B-A-G-C-C/D-D'-H F-E-B-A-G-C-C/D-D'-H-L F-E-B-A-G-C-C/D-D'-H-K-L F-E-B-A-G-C-C/D'-D F-E-B-A-G-C-C/D'-H F-E-B-A-G-C-C/D'-H-L F-E-B-A-G-C-C/D'-H-K-L F-E-B-A-G-C-C/D'-D'-H F-E-B-A-G-C-C/D'-D'-H-L F-E-B-A-G-C-C/D'-D'-'H-K-L F-E-B-A-G-C-C/D-C/D'-D F-E-B-A-G-C-C/D-C/D'-H F-E-B-A-G-C-C/D-C/D'-H-L F-E-B-A-G-C-C/D-C/D'-H-K-L F-E-B-A-G-C-C/D-C/D'-D'-H F-E-B-A-G-C-C/D-C/D'-D'-H-L F-E-B-A-G-C-C/D-C/D'-D'-H-K-L

Figure 17 B

E-B-A'

E-B-A-C-C/D-D E-B-A-C-C/D-H E-B-A-C-C/D-H-L E-B-A-C-C/D-H-K-L E-B-A-C-C/D-D'-H E-B-A-C-C/D-D'-H-L E-B-A-C-C/D-D'-H-K-L E-B-A-C-C/D'-D E-B-A-C-C/D'-H E-B-A-C-C/D'-H-L E-B-A-C-C/D'-H-K-L E-B-A-C-C/D'-D'-H E-B-A-C-C/D'-D'-H-L E-B-A-C-C/D'-D'-H-K-L E-B-A-C-C/D-C/D'-D E-B-A-C-C/D-C/D'-H E-B-A-C-C/D-C/D'-H-L E-B-A-C-C/D-C/D'-H-K-L E-B-A-C-C/D-C/D'-D'-H E-B-A-C-C/D-C/D'-D'-H-L E-B-A-C-C/D-C/D'-D'-H-K-L

E-B-A-G-C-C/D-D E-B-A-G-C-C/D-H E-B-A-G-C-C/D-H-L E-B-A-G-C-C/D-H-K-L E-B-A-G-C-C/D-D'-H E-B-A-G-C-C/D-D'-H-L E-B-A-G-C-C/D-D'-H-K-L E-B-A-G-C-C/D'-D E-B-A-G-C-C/D'-H E-B-A-G-C-C/D'-H-L E-B-A-G-C-C/D'-H-K-L E-B-A-G-C-C/D'-D'-H E-B-A-G-C-C/D'-D'-H-L E-B-A-G-C-C/D'-D'-H-K-L E-B-A-G-C-C/D-C/D'-D E-B-A-G-C-C/D-C/D'-H E-B-A-G-C-C/D-C/D'-H-L E-B-A-G-C-C/D-C/D'-H-K-L E-B-A-G-C-C/D-C/D'-D'-H E-B-A-G-C-C/D-C/D'-D'-H-L E-B-A-G-C-C/D-C/D'-D'-H-K-L WO 96/15812 PCT/US95/14974

AGC Ser 1	CAT His	CTT Leu	GTC Val	AAG Lys 5	TGT Cys	GCA Ala	GAG Glu	AAG Lys	GAG Glu 10	AAA Lys	ACT Thr	TTC Phe	TGT Cys	GTG Val 15	AAT Asn	4:
GGA Gly	GGC Gly	GAG Glu	TGC Cys 20	TTC Phe	ATG Met	GTG Val	AAA Lys	GAC Asp 25	CTT Leu	TCA Ser	AAT Asn	CCC Pro	TCA Ser 30	AGA Arg	TAC Tyr	96
TTG Leu	TGC Cys	AAG Lys 35	TGC Cys	CCA Pro	TAA Asn	GAG Glu	TTT Phe 40	ACT Thr	GGT Gly	GAT Asp	CGC Arg	TGC Cys 45	CAA Gln	AAC Asn	TAC Tyr	. 144
GTA Val	ATG Met 50	GCC Ala	AGC Ser	TTC Phe	Tyr	AGT Ser 55	ACG Thr	TCC Ser	ACT Thr	CCC Pro	TTT Phe 60	CTG Leu	TCT Ser	CTG Leu	CCT Pro	192
SAA Slu	TAG															198

PCT/US95/14974

AGC Ser 1	CAT His	CTT Leu	GTC Val	AAG Lys 5	TGT Cys	GCA Ala	GAG Glu	AAG Lys	GAG Glu 10	AAA Lys	ACT Thr	TTC Phe	TGT Cys	GTG Val 15	AAT Asn	4:	B
GGA Gly	GGC Gly	GAG Glu	TGC Cys 20	TTC Phe	ATG Met	GTG Val	AAA Lys	GAC Asp 25	CTT Leu	TCA Ser	AAT Asn	CCC Pro	TCA Ser 30	AGA Arg	TAC Tyr	9	6
TTG Leu	TGC Cys	AAG Lys 35	TGC Cys	CAA Gln	CCT Pro	GGA Gly	TTC Phe 40	ACT Thr	GGA Gly	GCG Ala	AGA Arg	TGT Cys 45	ACT Thr	GAG Glu	AAT Asn	14	4
GTG Val	CCC Pro 50	ATG Met	AAA Lys	GTC Val	CAA Gln	ACC Thr 55	CAA Gln	GAA Glu	AAA Lys	GCG Ala	GAG Glu 60	GAG Glu	CTC Leu	TAC Tyr	TAA	19	2

WO 96/15812

Ser 1	His	Leu	Val	Lys 5	Cys	Ala	GAG Glu	AAG Lys	GAG Glu 10	AAA Lys	ACT Thr	TTC Phe	TGT Cys	GTG Val 15	AAT Asn	48
GGA Gly	GGC Gly	GAG Glu	TGC Cys 20	TTC Phe	ATG Met	GTG Val	AAA Lys	GAC Asp 25	CTT Leu	TCA Ser	AAT Asn	CCC Pro	TCA Ser 30	AGA Arg	TAC Tyr	96
TTG Leu	TGC Cys	AAG Lys 35	TGC Cys	CCA Pro	AAT Asn	GAG Glu	TTT Phe 40	ACT Thr	GGT Gly	GAT Asp	CGC Arg	TGC Cys 45	CAA Gln	AAC Asn	TAC Tyr	144
GTA Val	ATG Met 50	GCC Ala	AGC Ser	TTC Phe	TAC Tyr	AAA Lys 55	GCG Ala	GAG Glu	GAG Glu	CTC Leu	TAC Tyr 60	TAA				183

WO 96/15812 PCT/US95/14974

AGC Ser 1	CAT His	CTT Leu	GTC Val	AAG Lys 5	TGT Cys	GCA Ala	GAG Glu	AAG Lys	GAG Glu 10	AAA Lys	ACT	Phe	Cys	Val 15	Asn	40
GGA Gly	GGC Gly	GAG Glu	TGC Cys 20	TTC Phe	ATG Met	GTG Val	AAA Lys	GAC Asp 25	CTT Leu	TCA Ser	AAT Asn	CCC Pro	TCA Ser 30	AGA Arg	TAC Tyr	96
TTG Leu	TGC Cys	AAG Lys 35	TGC Cys	CCA Pro	AAT Asn	GAG Glu	TTT Phe 40	ACT Thr	GGT Gly	GAT Asp	CGC Arg	TGC Cys 45	CAA Gln	AAC Asn	TAC Tyr	144
GTA Val	ATG Met 50	GCC Ala	AGC Ser	TTC Phe	TAC Tyr	AAG Lys 55	CAT His	CTT Leu	GGG Gly	ATT Ile	GAA Glu 60	TTT Phe	ATG Met	GAG Glu	AAA Lys	192
		GAG Glu			TAA											210

Ser 1	His	Leu	Val	Lys 5	Cys	'GCA Ala	GAG Glu	AAG Lys	GAG Glu 10	AAA Lys	ACT	TTC Phe	TGT Cys	GTG Val 15	AAT Asn	4
GGA Gly	GGC	GAG Glu	TGC Cys 20	TTC Phe	ATG Met	GTG Val	AAA Lys	GAC Asp 25	CTT Leu	TCA Ser	AAT Asn	CCC Pro	TCA Ser 30	AGA Arg	TAC Tyr	90
TTG Leu	TGC Cys	AAG Lys 35	TGC Cys	CAA Gln	CCT Pro	GGA Gly	TTC Phe 40	ACT Thr	GGA Gly	GCG Ala	AGA Arg	TGT Cys 45	ACT Thr	GAG Glu	AAT Asn	144
GTG Val	CCC Pro 50	ATG Met	AAA Lys	GTC Val	CAA Gln	ACC Thr 55	CAA Gln	GAA Glu	AAG Lys	TGC Cys	CCA Pro 60	AAT Asn	GAG Glu	TTT Phe	ACT Thr	192
GGT Gly 55	GAT Asp	CGC Arg	TGC Cys	CAA Gln	AAC Asn 70	TAC Tyr	GTA Val	ATG Met	GCC Ala	AGC Ser 75	TTC Phe	TAC Tyr	AGT Ser	ACG Thr	TCC Ser 80	240
ACT Thr	CCC Pro	TTT Phe	CTG Leu	TCT Ser 85	CTG Leu	CCT Pro	GAA Glu	TAG								. 267

AGC Ser 1	CAT His	CTT Leu	GTC Val	AAG Lys 5	TGT Cys	GCA Ala	GAG Glu	AAG Lys	GAG Glu 10	AAA Lys	ACT Thr	TTC Phe	TGT Cys	GTG Val 15	AAT Asn	48
GGA Gly	GGC Gly	GAG Glu	TGC Cys 20	TTC Phe	ATG Met	GTG Val	AAA Lys	GAC Asp 25	CTT Leu	TCA Ser	AAT Asn	CCC Pro	TCA Ser 30	AGA Arg	TAC Tyr	96
TTG Leu	TGC Cys	AAG Lys 35	TGC Cys	CAA Gln	CCT Pro	GGA Gly	TTC Phe 40	ACT Thr	GGA Gly	GCG Ala	AGA Arg	TGT Cys 45	ACT Thr	GAG Glu	AAT Asn	144
GTG Val	CCC Pro 50	ATG Met	AAA Lys	GTC Val	CAA Gln	ACC Thr 55	CAA Gln	GAA Glu	AAG Lys	TGC Cys	CCA Pro 60	AAT Asn	GAG Glu	TTT Phe	ACT Thr	192
GGT Gly 65	GAT Asp	CGC Arg	TGC Cys	CAA Gln	AAC Asn 70	TAC Tyr	GTA Val	ATG Met	GCC Ala	AGC Ser 75	TTC Phe	TAC Tyr	AAA Lys	GCG Ala	GAG Glu 80	240
	CTC Leu		TAA													252

Figure 24 A

001100000		
	TITT TITTITCTT NNTTTTTTT TGCCCTTATA CCTCTTCGCC	60
	CACT TCTTCCCCCT CCTCCTCCCA TAAACAACTC TCCTACCCCT	120
GCACCCCCAA TAAATAA	AATA AAAGGAGGAG GGCAAGGGGG GAGGAGGAGG AGTGGTGCTG	180
CGAGGGGAAG GAAAAGG	GGAG GCAGCGCGAG AAGAGCCGGG CAGAGTCCGA ACCGACAGCC	240
AGAAGCCCGC ACGCACC	TCG CACC ATG AGA TGG CGA CGC GCC CCG CGC CGC Met Arg Trp Arg Arg Ala Pro Arg Arg 1 5	291
10	C CCC CGG GCC CAG CGC CCC GGC TCC GCC GCC CGC y Pro Arg Ala Gln Arg Pro Gly Ser Ala Ala Arg 15 20 25	339
TCG TCG CCG CCG CTC Ser Ser Pro Pro Let	G CCG CTG CTG CCA CTA CTG CTG CTG GGG ACC u Pro Leu Pro Leu Leu Leu Leu Gly Thr Val Cys Leu Leu Thr Val	387
30	GGF II 09 35 40	
Ala Ala Leu Pro Pro		435
45	50 55	
GGG GCC TCG GTG TGC Gly Ala Ser Val Cys	TAC TCG TCC CCG CCC AGC GTG GGA TCG GTG CAG Tyr Ser Ser Pro Pro Ser Val Gly Ser Val Gln Ala Ser Pro Val Ser Val Gly Ser Val Gln	483
60	GGF II 08 70	
Glu Leu Val Gln Arg	GCC GCG GTG GTG ATC GAG GGA AAG GTG CAC CCG Ala Ala Val Val Ile Glu Gly Lys Val His Pro Trp Phe Val Val Ile Glu Gly Lys GGF II 04	531
75	80 85	

Figure 24 B

CAG Gln 90	CGG Arg	CGG Arg	CAG Gln	CAG Gln	GGG Gly 95	GCA Ala	CTC Ļeu	GAC Asp	AGG Arg	AAG Lys 100	GCG Ala	GCG Ala	GCG Ala	GCG Ala	GCG Ala 105	579
GGC Gly	GAG Glu	GCA Ala	GGG Gly	GCG Ala 110	TGG Trp	GGC Gly	GGC	GAT Asp	CGC Arg 115	GAG Glu	CCG Pro	CCA Pro	GCC Ala	GCG Ala 120	GGC Gly	627
CCA Pro	CGG Arg	GCG Ala	CTG Leu 125	GGG Gly	CCG Pro	CCC Pro	GCC Ala	GAG Glu 130	GAG Glu	CCG Pro	CTG Leu	CTC Leu	GCC Ala 135	GCC Ala	AAC Asn	675
GGG Gly	ACC Thr	GTG Val 140	CCC Pro	TCT Ser	TGG Trp	CCC Pro	ACC Thr 145	GCC Ala	CCG Pro	GTG Val	CCC Pro	AGC Ser 150	GCC Ala	GGC Gly	GAG Glu	723
CCC Pro	GGG Gly	GAG Glu	GAG Glu	GCG Ala	CCC Pro	TAT Tyr	Len CIG	GTG Val	LVS	Val	His	CAG Gln Glu 01	Val	Trp	Ala	771
	155					160					165					
Val	AAA Lys Lys	GCC Ala	GGG Gly	GGC Gly	TTG Leu	AAG Lys	AAG Lys	ASD	TCG Ser	CTG	CTC	ACC	GTG Val	CGC	CTG Leu	819
								Asp	Leu	GGF	Leu II	Aaa	Val	5	200	
170					175			Asp	Leu	GGF 180	II :	10	VAI		185	0.67
GGG	ጥኮሎ	TGG Trp Trp	Gly Gly	CAC His Pro	175 CCC Pro	GCC Ala	TTC Phe	CCC Pro	TCC	GGF 180 TGC Cys	II :	AGG Arg	CTC	AAG	185 GAG	867
GGG	ጥኮሎ	TYY	Gly Gly	CAC His	175 CCC Pro	GCC Ala	TTC Phe	CCC Pro	TCC	GGF 180 TGC Cys	II :	AGG Arg	CTC	AAG	185 GAG Glu	867
GGG Gly Gly	Thr	Trp	Gly Gly GGI TAC	CAC His Pro F II 190	175 CCC Pro Pro 03	GCC Ala Ala TTC	TTC Phe Phe ATG	CCC Pro Pro GAG Glu Glu	TCC Ser Val 195 CCC Pro	GGF 180 TGC Cys Xaa GAC Asp	GGG Gly Tyr GCC Ala	AGG Arg AAC	CTC Leu AGC	AAG Lys 200 ACC Thr Ser	185 GAG Glu	867 915

Figure 24 C

Ar	g Al	G CC a Pr 22	OAT	C GCG a Ala	TTO Phe	C CGA	A GCC J Ala 225	Ser	TTC Phe	CCC Pro	C CC	CT(Let 23(ه Gl	G ACC	GGC Gly	963
CGG	G AAG J Asi 23!	יו דיפו	C AAG u Ly:	G AAC s Lys	G GAG	GTC Val 240	. Ser	CGG	GTG Val	CTG Leu	TGC Cys 245	: Lys	G CGG	G TGO G Cys	GCC Ala	1011
Leu 250	LTC	CCC Pro	CAA Gli	A TTO	AAA Lys 255	Glu	ATG Met	AAA Lys	AGC Ser	CAG Gln 260	Glu	TCC Ser	GC: Ala	r GCA Ala	GGT Gly 265	1059
TCC Ser	AAA Lys	ner.	Val Val	CTT Leu Leu	Arg Arg	Cys	GAA Glu	ACC Thr	AGT Ser	TCT Ser	GAA Glu	TAC	Ser	TCT Ser	CTC Leu	1107
				270					175					180		
AGA Arg	TTC Phe	AAG Lys	TGG Trp 185	Pne	AAG Lys	AAT Asn	GGG Gly	AAT Asn 190	GAA Glu	TTG Leu	AAT Asn	CGA Arg	AAA Lys 195	Asn	AAA Lys	1155
CCA Pro	CAA Gln	AAT Asn 200	ATC Ile	AAG Lys	ATA Ile	CAA Gln	AAA Lys 205	AAG Lys	CCA Pro	GGG Gly	AAG Lys	TCA Ser 210	GAA Glu	CTT Leu	CGC Arg	1203
ATT Ile	AAC Asn	rys	Ala	TCA Ser Ser	Leu	Ala	Asp Asp	Ser Ser	Gly Gly	Glu	Tvr	Met	Cve	Tare	GTG Val	1251
	215					220	GG	F II	12		225					
ATC Ile 230	AGC Ser	AAA Lys	TTA Leu	GGA Gly	AAT Asn 235	GAC Asp	AGT (GCC (Ser .	GCC Ala 240	AAT Asn	ATC Ile	ACC Thr	ATC Ile	GTG Val 245	1299
GAA Glu	TCA Ser	AAC Asn	ATG	ACA Thr 250	TCT . Ser '	ACA (TCC 1 Ser 1	thr !	ACT (Thr (255	GGG . Gly '	ACA Thr	AGC Ser	CAT His	CTT Leu 260	GTA Val	1347

Figure 24 D

AAA Lys	TGT Cys	GCG Ala	GAG Glu 265	AAG Lys	GAG Glu	AAA Lys	ACT Thr	TTC Phe 270	TGT Cys	GTG Val	AAT Asn	GGA Gly	GGG Gly 275	GAG Glu	TGC Cys		1395
TTC Phe	ATG Met	GTG Val 280	AAA Lys	GAC Asp	CTT Leu	TCA Ser	AAC Asn 285	CCC Pro	TCG Ser	AGA Arg	TAC Tyr	TTG Leu 290	TGC Cys	AAG Lys	TGC Cys		1443
CCA Pro	AAT Asn 295	GAG Glu	TTT Phe	ACT Thr	GGT Gly	GAT Asp 300	CGC Arg	TGC Cys	CAA Gln	AAC Asn	TAC Tyr 305	GTA Val	ATG Met	GCC Ala	AGC Ser		1491
TTC Phe 400	TAC Tyr	AGT Ser	ACG Thr	TCC Ser	ACT Thr 405	CCC Pro	TTT Phe	CTG Leu	TCT Ser	CTG Leu 410	CCT Pro	GAA Glu			\		1530
TAG	SAGC	ATG (CTCAC	GTTG	ST GO	CTGC	rttc:	r TGʻ	TTGC:	IGCA	TCT	CCCC'	ICA (GATT	CCACC'	T	1590
AGA	CTAC	GAT (TGT(CTTAC	CC AC	GATC'	TAAT	A TTV	GACT	GCCT	CTG	CCTG'	rcg ·	CATG	AGAAC	A	1650
TTA	ACAA	AAG (CAAT!	rgta:	TT AC	CTTC	CTCT	G TT	CGCG	ACTA	GTT	GCT	CTG .	AGAT	ACTAA'	T	1710
															attga		1770
															TATTG		1830
															CCTGA		1890
															TTGAT		1950
	AATG:																2003

PGPRAQRP	61 VCYSSPPSVGSVQELAQRAAVVIEGKVHPQRRQQGALDRKAAAAAGEAGAWGGDREPPAA	121 GPRALGPPAEEPLLAANGTVPSWPTAPVPSAGEPGEEAPYLVKVHQVWAVKAGGLKKDSL 11-3 11-2 181 LTVRLGTWGHPAFPSCGRLKEDSRYIFFMEPDANSTSRAPAAFRASFPPLETGRNLKKEV	241 SRVLCKRCALPPQLKEMKSQESAAGSK 1 O OMSERKEGRGKGKKKKERGSGKKPESAAGSQSP 1 R K G D VP GP R	1 I-7, I III-13	328 4 II-12 5 5 8 R S 11-13 113 KVISKLGNDSASANITIVESNATSTS TITGMPASTEGAYVSSESPIRISVSTEGANTSSS	354 TTGTSHLVKÇAEKEKTFCVNGGECFMVKDLSNPSRYLCKCPNEFTGDRCQNYVMASFYST 173 A	9
GGFHBS5			GGFHBS5 GGFHFB1 GGFBPP5				

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Met 1	Arg	Trp	Arg	Arg 5	Ala	Pro	Arg	Arg	Ser 10	Gly	Arg	Pro	Gly	Pro 15	Arg
Ala	Gln	Arg	Pro 20	Gly	Ser	Ala	Ala	Arg 25	Ser	Ser	Pro	Pro	Leu 30	Pro	Leu
Leu	Pro	Leu 35	Leu	Leu	Leu	Leu	Gly 40	Thr	Ala	Ala	Leu	Ala 45	Pro	Gly	Ala
Ala	Ala 50	Gly	Asn	Glu	Ala	Ala 55	Pro	Ala	Gly	Ala	Ser 60	Val	Cys	Tyr	Ser
Ser 65	Pro	Pro	Ser	Val	Gly 70	Ser	Val	Gln	Glu	Leu 75	Ala	Gln	Arg	Ala	Ala 80
Val	Val	Ile	Glu	Gly 85	Lys	Val	His	Pro	Gln 90	Arg	Arg	Gln	Gln	Gly 95	Ala
Leu	Asp	Arg	Lys 100	Ala	Ala	Ala	Ala	Ala 105	Gly	Glu	Ala	Gly	Ala 110	Trp	Gly
Gly	Asp	Arg 115	Glu	Pro	Pro	Ala	Ala 120	Gly	Pro	Arg	Ala	Leu 125	Gly	Pro	Pro
Ala	Glu 130	Glu	Pro	Leu	Leu	Ala 135	Ala	Asn	Gly	Thr	Val 140	Pro	Ser	Trp	Pro
Thr 145	Ala	Pro	Val	Pro	Ser 150	Ala	Gly	Glu	Pro	Gly 155	Glu	Glu	Ala	Pro	Tyr 160
Leu	Val	Lys	Val	His 165	Gln	Val	Trp	Ala	Val 170	Lys	Ala	Gly	Gly	Leu 175	Lys
			180				Arg	185					130		
		195					Lys 200					203			
	210					215					220				
225					230		Thr			233					240
				245					250						Glu
			260					265					270		Cys
		275					280					203			Asn
	290					295					300				Gln
305	•				310					212	1				Ala 320
Asp	Ser	Gly	Glu	Tyr 325	Met	Сув	Lys	Val	11e 330	Ser	Lys	Lev	Gly	Asn 335	Asp

Figure 26 (cont.)

Ser Ala Ser Ala Asn Ile Thr Ile Val Glu Ser Asn Ala Thr Ser Thr 340 345 350

Ser Thr Thr Gly Thr Ser His Leu Val Lys Cys Ala Glu Lys Glu Lys 355 360 365

Thr Phe Cys Val Asn Gly Gly Glu Cys Phe Met Val Lys Asp Leu Ser 370 380

Asn Pro Ser Arg Tyr Leu Cys Lys Cys Pro Asn Glu Phe Thr Gly Asp 385 390 395 400

Arg Cys Gln Asn Tyr Val Met Ala Ser Phe Tyr Ser Thr Ser Thr Pro 405 410 415

Phe Leu Ser Leu Pro Glu * 420

TCT	AA AA Aa	CTI INT)	AC AC	SA GA	AC TO	FT AT 75 II 5	le Pi	ic Ai	et II	le Il	e II	le Va	al Le	eu Xa	aa A	sn 15	Ile	J J
CTT Leu	AAA Lys	CCG Pro	CTT Leu 20	TGG Trp	TCC Ser	TGA Xaa	TCT Ser	TGT Cys 25	AGG Arg	AAG Lys	TCA Ser	GAA Glu	CTT Leu 30	CGC Arg	ATT Ile	·		101
AGC Ser	AAA Lys	GCG Ala 35	TCA Ser	CTG Leu	GCT Ala	GAT Asp	TCT Ser 40	GGA Gly	GAA Glu	TAT Ser	ATG Met	TGC Cys 45	AAA Lys	GTG Val	ATC Ile	:		149
AGC Ser	AAA Lys 50	CTA Leu	GGA Gly	AAT Asn	GAC Asp	AGT Ser 55	GCC Ala	TCT Ser	GCC Ala	AAC Asn	ATC Ile 60	ACC Arg	ATT Ile	GTG Val	GAC Glu	3 1		197
TCA Ser 65	AAC Asn	GGT Gly	AAG Lys	AGA Arg	TGC Cys 70	CTA Leu	CTG Leu	CGT Arg	GCT Ala	ATT Ile 75	TCT Ser	CAG Gln	TCT Ser	CTA Leu	AGA Arg 80	a A		245
GGA Gly	GTG Val	ATC Ile	AAG Lys	GTA Val 85	TGT Cys	GGT Gly	CAC His	ACT Thr	TGA Xaa 90	ATC Ile	ACG Thr	CAG Gln	GTG Val	TGT Cys 95	GAI Glu	A 1		293
ATC Ile	TCA Ser	TTG Cys	TGA Xaa 100	ACA Thr	AAT Asn	AAA Lys	AAT Asn	CAT His 105	GAA Glu	AGG Arg	AAA Lys	ACT Thr	CTA Leu 110	TGT Cys	TTO	3 1		341
AAA Lys	TAT Tyr	CTT Leu 115	ATG Met	GGT Gly	CCT Pro	CCT Pro	GTA Val 120	AAG Lys	CTC Leu	TTC Phe	ACT Thr	CCA Pro 125	TAA Xaa	GGT Gly	GAI Gl	A 1		389
Ile	GAC Asp	Leu	Lys	Tyr	ATA Ile	TAG Xaa 135	ATT Ile	ATT Ile	T									417

INTERNATIONAL SEARCH REPORT

International application No. PCT/US95/14974

									
A. CL IPC(6)	ASSIFICATION OF SUBJECT MATTER :A61K 48/00, 38/18								
US CL	:424/93.21; 514/12, 44 to International Patent Classification (IPC) or to	hath setional	olegnification and IDC						
	CLDS SEARCHED	Cour national	CHISTICAGON AND IFC	•					
Minimum	documentation searched (classification system follows)	owed by clas	sification symbols)						
U.S . :	424/93.21; 514/12, 44			!					
Document	ation searched other than minimum documentation to	o the extent t	hat such documents are include	d in the fields searched					
Electronic	data base consulted during the international search	n (name of da	ita base and, where practicable	. scarch terms used)					
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C. DOC	CUMENTS CONSIDERED TO BE RELEVANT	r							
Category*	Citation of document, with indication, where	e appropriate	, of the relevant passages	Relevant to claim No.					
×	FASEB Journal, Volume 8, Numbers 4-5, issued 1994, 36-37 Danilenko et al, "Neu Differentiation Factor (NDEF) Accelerates Epidermal Migration and Differentiation in Excisional Wounds", page A535, see entire abstract.								
A	BioEssays, Volume 15, Number 12, issued December 1993, Peles et al, "Neu and its Ligands: From an Oncogene to Neural Factors", pages 815-824, see entire document.								
A, P	US, A, 5,367,060 (VANDLEN E see column 37, line 54 to column			1-2, 7-32, 35- 37					
<u> </u>									
	r documents are listed in the continuation of Box (See patent family annex.						
A* docum	ment defining the general state of the art which is not considered		inter document published after the inter late and not in conflict with the applicat principle or theory underlying the inves	ion but cited to understand the					
	of particular relevance r document published on or after the international filling date	•x•	locument of particular relevance; the	claimed invention cannot be					
." docum	ment which may throw doubts on priority claim(s) or which is to establish the publication dots of another citation or other	•	considered movel or cannot be considered when the document is taken alone	d to involve an inventive step					
specia	d reason (as apocified) most referring to an oral disclosure, use, exhibition or other	G	ocument of particular relevance; the considered to involve an inventive a combined with one or more other such (top whos the document in					
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the pr	tual completion of the international search	-	ailing of the initernational scan	•					
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INTERNATIONAL SEARCH REPORT

International application No. PCT/US95/14974

C (Continua	ation). DOCUMENTS CONSIDERED TO BE RELEVANT	
Category*	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
A	J. P. BROCKES et al, "The Neuron as a Source of Mitogen: Its Influence on the Proliferation of Glial and Non-neural Cells," from "Development in the Nervous System", published 1980, Garrod and Feldman, (CA), pages 309-327, see entire document.	1-2, 7-32, 35-37
A, P	US, A, 5,399,346 (ANDERSON ET AL) 21 March 1995, see entire document.	3-6, 33-34
A	Proc. Natl. Acad. Sci. USA, Volume 89, issued April 1992, Yao et al, "Expression of Human Factor IX in Mice after Injection of Genetically Modified Myoblasts", pages 3357-3361, see entire document.	3-6, 33-34
А, Р	SPECIAL NEWS REPORT, Volume 269, issued 25 August 1995, Marshall, "Gene Therapy's Growing Pains", pages 1050-1055, see entire document.	3-6, 33-34
A	US, A, 5,082,670 (GAGE ET AL) issued 21 January 1992, see entire document.	3-6, 33-34

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B. FIELDS SEARCHED

Electronic data bases consulted (Name of data base and where practicable terms used):

APS/USPAT, MEDLINE, BIOSIS, WORLD PATENT INDEX

Search terms: neuregulin#, heregulin#, glial growth factor#, acetylcholine receptor inducing activity, ARIA, neu differentiation factor#

BOX II. OBSERVATIONS WHERE UNITY OF INVENTION WAS LACKING

This ISA found multiple inventions as follows:

This application contains the following inventions or groups of inventions which are not so linked as to form a single inventive concept under PCT Rule 13.1. In order for all inventions to be examined, the appropriate additional examination fees must be paid.

Group I, claim(s)1-2, 7-32, and 35-37, drawn to a method of affecting cellular communication in a vertebrate by administering a neurogulin protein.

Group II, claim(s) 3-6 and 33-34, drawn to a method of affecting cellular communication in a vertebrate by adminstering neurogulin-producing cells or neurogulin DNA (gene therapy).

The inventions listed as Groups I-II do not relate to a single inventive concept under PCT Rule 13.1 because, under PCT Rule 13.2, they lack the same or corresponding special technical features for the following reasons:

The special technical feature of Group I is the administration of neuregulin protein to vertebrates. The special technical feature of Group II is the administration of neuregulin DNA either directly (in vivo gene therapy) or after introduction of DNA into a cell (ex vivo gene therapy). The method of Group II does not require the administration of neuregulin protein and the method of Group I does not require the administration of neuregulin DNA. Furthermore, the method steps for administration of protein and DNA are different. Therefore, both the reagents used and the process steps of Groups I and II are distinct from one another and the special technical features of each method distinguish the two groups from one another. Accordingly, the claims are not so linked by a special technical feature within the meaning of PCT Rule13.2 so as to form a single inventive concept.

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